STIC-Biotech/ChemLib

Li, Ruixiang

ଅFr m: Sent:

Wednesday, March 19, 2003 10:56 AM STIC-Biotech/ChemLib

T:

Subject:

Sequence search of Application NO: 09/727,739

Please do a standard search on:

(i) SEQ ID NOS: 15 and 18 against the interference amino acid databases;

(ii) SEQ ID NOS: 17 and 19 against both commercial and interference amino acid databases.

Thank you very much!

Ruixiang Li GAU 1646 CM1 10E18 Mail Box 10D19 306-0282

TYPE OF SEARCH:

Searcher:___ Phone:____

Point of Contact P. Sheppard

Telephone number: (703) 308-4499

Location:_

Date Picked Up:__

Date Completed: 3/25/03

Searcher Prep/Review:____

Clerical:____

Online time:

NA Sequences:_____ AA Sequences:_____ Structures: Bibliographic: Litigation:

Full text: Patent Family:

Other:_____

VENDOR/COST (where applic.) STN:_____ DIALOG:_____ Questel/Orbit:_____ DRLink:_____ Lexis/Nexis:_____ Sequence Sys.:_____ WWW/Internet:____ Other (specify):_____

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10666 No. US20020013456Alth
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IG DATE: 18-Jan-2001
IFICATION: <Unknown>
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OPERATING SYSTEM: PC-
SOFTWARE: Patentin Re-
CURRENT APPLICATION DATA:
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2: /cgn2_6/ptodata/1/pubpaa/DS06_NEW_PUB.pep
3: /cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep
4: /cgn2_6/ptodata/1/pubpaa/US07_NEW_PUB.pep
5: /cgn2_6/ptodata/1/pubpaa/US07_NEW_PUB.pep
6: /cgn2_6/ptodata/1/pubpaa/US07_NEW_PUB.pep
7: /cgn2_6/ptodata/1/pubpaa/US07_NEW_PUB.pep
8: /cgn2_6/ptodata/1/pubpaa/US09_NEW_PUB.pep
9: /cgn2_6/ptodata/1/pubpaa/US09_NEW_PUB.pep
10: /cgn2_6/ptodata/1/pubpaa/US09_NEW_PUB.pep
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12: /cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep
13: /cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep
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US-10-062-375-3
Sequence 3, Application US/10062375
Fatent No. US20020133000A1
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Sutcliffe, Gregor J.
de Lecea, Luis
Siggins, George R.
Henriksen, Steven J.
TITLE OF INVENTION: CORTISTATIN: NEUROPEPTIDES,
COMPOSITIONS AND METHODS
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: THE SCRIPPS RESEARCH INSTITUTE
STREET: 10666 No. US20020133000A1th forrey Pines R
CITY: La Jolla
STATE: Callifornia
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/062,375
FILING DATE: 30-Jan-2002
CLASSIFICATION: CURROWN>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/857,389
FILING DATE: CURROWN>
ATTORNEY/AGENT INFORMATION:
NAME: Schmonsees, William
REGISTRATION NUMBER: 31,796
REFERENCE/DOCKET NUMBER: 22908-0002
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFAX: (415) 324-7041
TELEFAX: (415) 324-0638
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larity 38.3%; Pred. No. 7.9e-11;
Conservative 17; Mismatches 30;
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Best Local Similarity 38.3%; Pred. No. 7.90
Matches 46; Conservative 17; Mismatches
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SEQUENCE CHARACTERISTICS:

LENGTH: 110 amino acids

TYPE: amino acid

TOPOLOGY: 11near

MOLECULE TYPE: protein

FRAGMENT TYPE: C-terminal

SEQUENCE DESCRIPTION: SEQ ID NO: 3:

US-10-062-375-3
TOPOLOGY: linear

MOLECULE TYPE: protein

FRAGMENT TYPE: C-terminal

SEQUENCE DESCRIPTION: SEQ ID NO: 3

US-09-766-396-3
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an amino acid sequence of
MWPsp-MWPmp20-(His)6-EGF-TEV-Somatostatin
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Pred. No. 0.00015;
18; Mismatches 35;
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CURRENT FILING DATE: 1999-03-
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CORRESPON.

STREET: 10666 ...

STREET: 20166 ...

CITY: La Jolla

STATE: California

COUNTRY: US

COUNTRY: US

CONFORT: IBM PC COMPATIBLE

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC COMPATIBLE

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PATENTIN Release #1.0, Version #1.25

CURRENT APPLICATION NUMBER: US/09/766,396

FILING DATE: 18-Jan-2001

CLASSIFICATION NUMBER: US/09/766,396

FILING DATE: 18-Jan-2001

CLASSIFICATION NUMBER: 08/857,389

FILING DATE: CURKNOWN-

ATTORNEY/AGENT INFORMATION:

NAME: SCHMONSEES, William

REGISTRATION NUMBER: 22908-0002

TELECOMMUNICATION INFORMATION:

TELERNOEN: (415) 324-7041

TELERNOEN: (415) 324-7041

TELERNOEN: (415) 324-0638

INFORMATION FOR SEQ ID NO: 2:

SEQUENCE CHARACTERISTICS:

TELENGTH: 112 amino acids

TYPE: amino acids

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Sutcliffe, Gregor J.

de Lecea, Luis
Siggins, George R.
Henriksen, Steven J.

TITLE OF INVENTION: CORTISTATIN: NEUROPEPTIDES,
COMPOSITIONS AND METHODS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: THE SCRIPPS RESEARCH INSTITUTE
STREET: 10666 No. US20020013456A1th Torrey
                                                                                                                      Artificial
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                                                                                                                                                                                                                                                                                                                CTHER INFORMATION: Description of CTHER INFORMATION: protein US-101-487-53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 2, Application US/09766396
Patent No. US20020013456A1
GENERAL INFORMATION:
APPLICANT: Sutcliffe, Gregor
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                                                                     Sequence
                                                                                                                                                                                                               Ouery Match
Best Local Similarity 48.4%;
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         rOPOLOGY: linear;
MOLECULE TYPE: protein;
SEQUENCE DESCRIPTION: S
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                                        TYPE: PRT ORGANISM: Artificial FEATURE:
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SEQ ID NO 53
LENGTH: 200
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US-09-766-396-2
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Gaps
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Pred. No. 0.014;
); Mismatches
                                                                                                                                                                                                                                                                                                                                                             NUMBER: US/08/857,389
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                                                                                                                                                                                                                                                                                          Patentin Release #1.
                                                                                                                                                                                                                                                                                                              UMBER: US/10/062,
30-Jan-2002
                                                                                                                      STATIN:
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EWASQDSSSTAFEGGTPELSKRQ
                                                                                                                                                                                                                                                                                                                                                                                               William
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0133000A1
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25.2%;
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                                                                                                                                                                                        No.
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Siggins, (
                                                                                                                                                                               THE
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                                                                                                                                  INVENTION:
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020013456A1
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POLKAKEDLERSVDNLPPRER
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SYSTEM
                                                 (FICATION:
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Best Local Similarity
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                                                                                                                  ATTORNEY
                                                                                                                                                                                                                                                     SEQUENCE
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GENERAL
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                                                                                                 Road,
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US-10-062-375-6
Sequence 6, Application US/10062375
Sequence 6, Application US/10062375
Sequence No. US20020133000A1
GENERAL INFORMATION:
CENERAL INFORMATION:
CENERAL INFORMATION:
CORPOSITIONS AND METHODS
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
STREET: 10666 No. US20020133000A1th Torrey Pines R
CITY: La Jolla
STATE: California
                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/766,396
FILING DATE: 18-Jan-2001
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/857,389
FILING DATE: <Unknown>
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                         NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: THE SCRIPPS RESEARCH INSTITUTE
STREET: 10666 No. US20020013456Alth Torrey Pine
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Henriksen, Steven J.
INVENTION: CORTISTATIN: NEUROPEPTIDES
COMPOSITIONS AND METHODS
7 SEQUENCES: 26
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Best Local Similarity 44.4%; Pred. No. 0.029;
Matches 16; Conservative 5; Mismatches 1
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NAME: Schmonsees, William
REGISTRATION NUMBER: 31,796
REFERENCE/DOCKET NUMBER: 22908-0002
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 324-7041
TELEFAX: (415) 324-0638
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SEQUENCE CHARACTERISTICS:
LENGTH: 85 amino acids
TYPE: amino acid
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MEDIUM TYPE: Flopp
COMPUTER: IBM PC C
                                                                                                              CITY: La Jolla
STATE: California
COUNTRY: US
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                 TITLE OF
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8-Jan-2001
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DATA
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Gaps

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Indels

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Mismatches

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Score 80.5; DB 12; Pred. No. 0.07;

Length 105;

WRPRKVKGPQL-KAK 81

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2002-02-01
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020177165A1
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Best Local Similarity
      US-10-062-375-26
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Best Local Similarity 27.7%; Pred. No. 0.07;
Matches 26; Conservative 12; Mismatches 37; Indels 19
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COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: TBM FC compatible
OCNEUTER: TBM FC compatible
OFFENTING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PALENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/062,375
FILING DATE: 30-Jan-2002
CLASSIFICATION NUMBER: US/08/857,389
FILING DATE: CUNKNOWN>
APPLICATION NUMBER: US/08/857,389
FILING DATE: CUNKNOWN>
APPLICATION NUMBER: 31,796
REPERENCE/DOCKET NUMBER: 31,796
REFERENCE/DOCKET NUMBER: 3
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CORRESPONDENCE ADDRESS:
ADDRESSEE: THE SCRIPPS RESEARCH INSTITUTE
STREET: 10666 No. US20020133000A1th Torrey Pine
CITY: La Jolla
STATE: California
COUNTRY: US
COUNTRY: US
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US-10-062-375-26
; Sequence 26, Application US/10062375
; Patent No. US20020133000A1
; GENERAL INFORMATION:
; APPLICANT: Sutcliffe, Gregor J.
de Lecea, Luis
Siggins, George R.
Henriksen, Steven J.
Henriksen, Steven J.
COMPOSITIONS AND METHODS
                                                                                                                            ANTI-SENSE: NO FRAGMENT TYPE: Internal SEQUENCE DESCRIPTION: SEQ ID NO: 26: US-09-766-396-26
LENGTH: 105 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: NO
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R FILING DATE: 1997-11-25

R APPLICATION NUMBER: 60/069694

R FILING DATE: 1998-02-09

R APPLICATION NUMBER: 60/074092

R FILING DATE: 1998-02-09

R APPLICATION NUMBER: 60/074092

R FILING DATE: 1998-02-09

R APPLICATION NUMBER: 60/09294

R APPLICATION NUMBER: 60/09598

R APPLICATION NUMBER: 60/09901

R FILING DATE: 1998-04-08

R APPLICATION NUMBER: 60/09901

R FILING DATE: 1998-09-10

R APPLICATION NUMBER: 60/09901

R FILING DATE: 1998-09-10

R APPLICATION NUMBER: 60/09901

R FILING DATE: 1998-09-10

R APPLICATION NUMBER: 60/109601

R FILING DATE: 1998-09-10

R APPLICATION NUMBER: 60/109304

R APPLICATION NUMBER: 60/109396

R APPLICATION NUMBER: 60/109306

R APPLICATION NUMBER: 60/109495

R APPLICATION NUMBER: 09/136804

R APPLICATION NUMBER: 09/13
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/US98/14552 PCT/US99/05028 PCT/US98/25190 /US98/19330 PCT/US98/25108 /US98/19093 /US98/24855 /US99/20111 /US99/20594 APPLICATION NUMBER: PCT/US99/21547 PCT/US98/18824 /US98/19437 PCT/US99/12252 7380138 7808689 767609 802706 866028 886342 665350 709238 872035 380139 548815 870574 380137 403297 423844 PCT (* 09-7 : PCT/ 09-08 91-60 06-01 09 - 1509/ -60 PRIOR

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SER: 60/0
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SER: 60/0
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3R: 60/0
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                                                                 SQASAGPLIGEEA 120
               Gaps
                                       --WRPRKVKGPQL-KAK 81
              19;
               Indels
                                                              61 LLSGATATAALPLEGGPTGRDSEHMQEAAGIRKSSLLTFLAWWFEWT
              37;
0.11;
                                                                                       Pred. No. 0.1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTATION NUMBER: US/10/028,072
CURRENT APPLICATION NUMBER: 00/049911
PRIOR APPLICATION NUMBER: 60/049911
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-06-18
PRIOR FILING DATE: 1997-08-26
PRIOR FILING DATE: 1997-09-17
PRIOR PPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR PPLICATION NUMBER: 60/059115
PRIOR PPLICATION NUMBER: 60/059117
PRIOR PPLICATION NUMBER: 60/059117
PRIOR FILING DATE: 1997-09-17
PRIOR PPLICATION NUMBER: 60/059184
PRIOR PPLICATION NUMBER: 60/059184
PRIOR PPLICATION NUMBER: 60/059184
PRIOR PPLICATION NUMBER: 60/059184
PRIOR PPLICATION NUMBER: 60/059186
PRIOR PLING DATE: 1997-09-19
PRIOR PLING DATE: 1997-09-19
PRIOR PLING DATE: 1997-09-19
PRIOR PLING DATE: 1997-09-19
PRIOR FILING DATE: 1997-09-19
PRIOR PLING DATE: 1997-09-19
PRIOR FILING DATE: 1997-09-19
PRIOR FILING DATE: 1997-10-17
PRIOR PPLICATION NUMBER: 60/062287
PRIOR PLING DATE: 1997-10-17
                                    37 LLORALA-AALPHRSGVSERWRTFYPNCPCLR--
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US-10-028-072-380
; Sequence 380, Application US/10028072
; Publication No. US20030004311A1
GENERAL INFORMATION:
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7-10-24
ER: 60/063045
7-10-24
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            12;
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Beresini, Maureen
DeForge, Laura
Desnoyers, Luc
Filvaroff, Ellen
Gao, Wei-Qiang
Gerritsen, Mary E.
Goddard, Audrey
Godowski, Paul J.
Gurney, Austin L.
Sherwood, Steven
Smith, Victoria
Stewart, Timothy A.
Tumas, Daniel
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T: Wood, William
T: Zhang
INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PPLICATION NUMBER: 60/(PPLICATION NUMBER: 60/(TATE) 1997-10-17
Similarity 27.7%; 26; Conservative
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Best Local
Matches 2
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                     -WRPRKVKGPQL-KAK 81
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Pred. No. 0
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                     -AALPHRSGVSERWRTFYPNCPCLR
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. US20030022239A1
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Similarity 27.7%;
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US-10-121-049-380
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US-10-121-049-380
                     LLQRALA
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LENGTH: 155
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Best Local Similarity 27.7%; Pred. No. 0.11;
Matches 26; Conservative 12; Mismatches
PRIOR APPLICATION NUMBER: 60/081918
PRIOR FILING DATE: 1998-04-15
PRIOR FILING DATE: 1998-04-24
PRIOR FILING DATE: 1998-04-28
PRIOR PELICATION NUMBER: 60/08332
PRIOR FILING DATE: 1998-04-28
PRIOR PELICATION NUMBER: 60/084627
PRIOR APPLICATION NUMBER: 60/084637
PRIOR APPLICATION NUMBER: 60/084637
PRIOR PELING DATE: 1998-05-07
PRIOR PELING DATE: 1998-05-12
PRIOR PELING DATE: 1998-05-13
PRIOR PELING DATE: 1998-05-14
PRIOR PELING DATE: 1998-05-15
PRIOR PELING DATE: 1998-05-17
PRIOR PELING DATE: 1998-05-18
PRIOR PELING DATE: 1998-05-19
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US-10-140-470-380
US-10-140-470-380
Sequence 380, Application US/10140470
Publication No. US20030022331A1
GENERAL INFORMATION:
APPLICANT: BACF, Kevin P.
APPLICANT: Desnoyers, Luce
APPLICANT: Gerrisen, Mary E.
APPLICANT: Gerrisen, Mary E.
APPLICANT: Godowski, Paul J.
APPLICANT: Godowski, Paul J.
APPLICANT: Godowski, Paul J.
APPLICANT: Gurney, Austin L.
APPLICANT: Sherwood, Steven
APPLICANT: Sherwood, Steven
APPLICANT: When I wood, William
APPLICANT: When I wood, William
APPLICANT: ACTOR I Sherwood, Steven
APPLICANT: Tumas, Daniel
APPLICANT: ACTOR I Sherwood, Steven
APPLICANT: ACTOR I Sherwood, Steven
APPLICANT: ACTOR I Sherwood, Steven
APPLICANT: ACTOR INCONING THE SAME
TITLE OF INVENTION: ACTOR ENCODING THE SAME
TITLE OF INVENTION NUMBER: US/10/140.470
CURRENT FILING DATE: 2002-05-06
PHORY HISS
TYPE: PRT
                                                                                                                                                 IDES AND NUCLEIC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    . 37; Indels
APPLICANT: Smith, Victoria
APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Watanabe, Colin K
APPLICANT: Wood, William
APPLICANT: Wood, William
APPLICANT: Wood, William
APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDE
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3330R1C54
CURRENT APPLICATION NUMBER: US/10/123,904
CURRENT FILING DATE: 2002-04-16
Prior Application removed - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 550
SEQ ID NO 380
LENGTH: 155
TYPE: PRT
ORGANISM: Homo Sapien
SS-10-123-904-380
                                                                                                                                                                                                                                                                                                                                                                                                                               DB 9; Leng1
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Best Local Similarity 27.7%; Pred. No. 0.11
Matches 26; Conservative 12; Mismatches
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US-10-140-470-380
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AND METHODS
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Mismatches
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                     US-09-120-
US-09-144-
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5212156-1
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Pred. No.
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Release #1.0
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COMPOSITIONS
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APPLICATION NUMBER: US/08/648
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NUMBER: 51
                                                                                                                                                                                                                 Gregor
Luis
                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Fitting, Thomas
REGISTRATION NUMBER: 34,16
REFERENCE/DOCKET NUMBER: 5
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38.3%;
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No. 60748
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(619) 554-63
(619) 554-63
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MEDIUM TYPE: Floppy
COMPUTER: IBM PC COM
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APPLICANT: Sutcliffe,
APPLICANT: de Lecea, I
TITLE OF INVENTION: CC
TITLE OF INVENTION: CC
NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
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California
Y: US
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MOLECULE TYPE:
FRAGMENT TYPE:
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CITY: I
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    /cgn2_6/ptodata/2/laa/6A_COMB.pep:*
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US-08-488-159-1

US-08-465-764-1

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                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/648,322
FILING DATE:
CLASSIFICATION: 435
                                                                                                                                                                            APPLICANT: Sutcliffe, Gregor J.
APPLICANT: de Lecea, Luis
TITLE OF INVENTION: CORTISTATIN: NEUROPEPTIDES,
TITLE OF INVENTION: COMPOSITIONS AND METHODS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 3, Application US/09001472
; Patent No. 6232100
; GENERAL INFORMATION:
    APPLICANT: OLSEN, HENRIK S.
    APPLICANT: RUBEN, STEVEN M.
    TITLE OF INVENTION: CORTISTATIN. POLYPEPTIDES
    NUMBER OF SEQUENCES: 11
    CORRESPONDENCE ADDRESS:
    ADDRESSEE: STERNE, KESSLER, GOLDSTEIN & FOX P
    STREET: 1100 NEW YORK AVENUE, SUITE 600
    CITY: WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 87; DB 3;
Pred. No. 0.0018;
); Mismatches 4.
                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
ADDRESSEE: THE SCRIPPS RESEARCH INSTITUTE
STREET: 10666 No. 6074872th Torrey Pines R
CITY: La Jolla
STATE: California
COUNTRY: US
ZIP: 92037
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NAME: Fitting, Thomas
REGISTRATION NUMBER: 34,163
REFERENCE/DOCKET NUMBER: 519.0
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 554-2937
TELEFAX: (619) 554-6312.
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 112 amino acids
TYPE: amino acid
                                                                                                      Sequence 2, Application US/08648322
Sequence 2, Application US/08648322
Patent No. 6074872
GENERAL INFORMATION:
APPLICANT: Sutcliffe, Gregor J.
APPLICANT: de Lecea, Luis
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US-08-648-322-2
RWRPRKVKGPQLKAKE-
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US-09-001-472-3
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AND METHODS
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Mismatches
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                                                    31-DEC-1996
ON DATA:
                                                                                                             ER: US 60/037,386
-FEB-1997
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/ENTION: CORTISTATIN:
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6074872
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Release
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Sutcliffe, Grey-
                                                                                                                                        INFORMATION:
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amino acid
             DATA:
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30; Conservative
                                    Herewith
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                                                                                                                                                                  NUMBER:
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CORRESPONDENCE ADDRESS
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PatentIn
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                          APPLICATION NUMBER:
                                                                                                                 ION NUMBER:
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California
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             APPLICATION
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O. 6074872th Torrey Pines Road,
                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPAtible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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AND METHODS
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                                                                             REVARROEGAPPOQSARRDRMPCRNFFWKTFSSC
                                                            RERKAGCKNFYWKGFTSC
      LLORALA-AALPHRSGVSERWRTFYPNCPCLR
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--- 519.0
                                                                                                                                                                                                       APPLICANT: Sutcliffe, Gregor J. APPLICANT: de Lecea, Luis TITLE OF INVENTION: CORTISTATIN: TITLE OF INVENTION: COMPOSITIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sutcliffe, Gregor J. de Lecea, Luis TENTION: CORTISTATIN: TENTION:
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                                                                                                                                                                 US/08648322
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6074872
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31.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: Fitting, Thomas
REGISTRATION NUMBER: 34,
REFERENCE/DOCKET NUMBER:
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Patent No. 6074872
GENERAL INFORMATION:
APPLICANT: Sutclif
                                                             EDLERSVDNLPP-
                                                                                                                                                                           Patent No. 6074872
GENERAL INFORMATION
APPLICANT: Sutcl
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US-08-648-322-5
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/001,472
                                                                                                                                                                                                                                                                                                                                                                 Sequence 2, Application US/09001472; Patent No. 6232100; GENERAL INFORMATION:
APPLICANT: OLSEN, HENRIK S.
APPLICANT: RUBEN, STEVEN M.
TITLE OF INVENTION: CORTISTATIN POLYPEPTIDES
NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSEE: STERNE, KESSLER, GOLDSTEIN & FOX P
STREET: 1100 NEW YORK AVENUE, SUITE 600
CITY: WASHINGTON
                                                                                                                                                                                                    . No. 0.0041;
Mismatches 1.
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Pred. No. 0
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Pred. No.
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Herewith
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1488
REGISTRATION NUMBER: 34,163
REFERENCE/DOCKET NUMBER: 519.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 554-2937
TELEFAX: (619) 554-6312
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 85 amino acids
                                                                                                                                                                                                                                       5,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: STEFFE, ERIC K.
REGISTRATION NUMBER: 36,688
REFERENCE/DOCKET NUMBER: 141
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                      13.98;
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illarity 27.7%;
Conservative 1
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202) 371-2540
SEQ ID NO: 2
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16; Conservative
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INFORMATION FOR SEQ ID NO
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26; Conser
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FRAGMENT TYPE:
US-08-648-322-6
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-09-001-472-2
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STATE:
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US-09-001-472
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                                                                                                                                                                                Score 7
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                                                                                        TOPOLOGY
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Best Local
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US-08-648-322-11

Sequence 11, Application US/08648322

Patent No. 6074872

GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: Ge Lecea, Luis

TITLE OF INVENTION: COMPOSITIONS AND METHODS

NUMBER OF SEQUENCES: 24

CORRESPONDENCE ADDRESS:

ADDRESSEE: THE SCRIPPS RESEARCH INSTITUTE

STREET: 10666 No. 6074872th Torrey Pines Road, TPC-8

CITY: La Jolla

STATE: California

COUNTRY: US

ZIP: 92037
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
13.1%; Score 78; DB 3; Length 2
Best Local Similarity 51.9%; Pred. No. 0.004;
Matches 14; Conservative 2; Mismatches 11; Indel
                                                                                                                   COMPUTER FACABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/648,322
FILING DATE:
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Fitting, Thomas
REGISTRATION NUMBER: 519.0
TELEPHONE: (619) 554-2937
TELEPHONE: (619) 554-6312
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 29 amino acids
TYPE: amino acids
TYPE: amino acids
TOPOLOGY: linear
MOLECULE TYPE: Protein
FRAGMENT TYPE: C-terminal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/648,322
FILING DATE:
                  THE SCRIPPS RESEARCH INSTITUTE 566 NO. 6074872th Torrey Pines 1
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ATTORNEY/AGENT INFORMATION:
NAME: Fitting, Thomas
REGISTRATION NUMBER: 34,163
REFERENCE/DOCKET NUMBER: 519.0
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 554-2937
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS
                                                      La Jolla
California
                                   10666
                                                                    STATE: Cali
COUNTRY: US
ZIP: 92037
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              Release #1.0, Version
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Khoury-Christianson, Anastasia
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                                            BER: US/07/977,628A
7-NOV-1992
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NT INFORMATION:
5405597nan, Kevin
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                                 LICATION DATA
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APPLICATION DATA
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CHARACTERISTIC
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REFERENCE/DOCKET NU
TELECOMMUNICATION INF
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Patent No
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US-07-628A-1
Sequence 1, Application US/07977628A
Fatent No. 5405597
GENERAL INFORMATION:
APPLICANT: Dean, Richard T
APPLICANT: Lister-James, John
TITLE OF INVENTION: Technetium-99m Labeled Somatostatin-
TITLE OF INVENTION: Derived Peptides for Imaging
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Allegretti & Witcoff, Ltd.
STREET: 10 South Wacker Drive, Suite 3000
CITY: Chicago
STATE: Illinois
COUNTRY: USA
GENERAL INFORMATION:

APPLICANT: POTTER, ANDREW A.

APPLICANT: BEDMOND:

APPLICANT: HUGHES, HUW P.A.

TITLE OF INVENTION: ENHANCED INMUNOGENICITY USING LEU
TITLE OF INVENTION: CHIMERAS

NUMBER OF SEQUENCES: 15

CORRESPONDENCE ADDRESS: 15

COMPITY: PALO ALTO

STREET: 285 HAMILTON AVENUE, SUITE 200

CITY: PALO ALTO

STREET: 285 HAMILTON AVENUE, SUITE 200

CITY: PALO ALTO

STREET: 285 HAMILTON AVENUE, SUITE 200

CITY: PALO ALTO

STREET: 285 HAMILTON AVENUE, SUITE 200

CITY: PALO ALTO

STREET: 285 HAMILTON AVENUE, SUITE 200

CITY: PALO ALTO

STREET: 285 HAMILTON AVENUE, SUITE 200

CITY: PALO ALTO

STREET: 285 HAMILTON AVENUE, SUITE 200

COMPITR: PALO ALTO

STREET: ABOUT STREET: BODDY 418K

COMPUTER READBALE FORM: BC-DOS/MS-DOS

SOFTWARE: BAR PC COMPATION

APPLICATION NUMBER: US. 087/960,932

FILING DATE: 11-0CT-1995

CLASSIFICATION NUMBER: 33,208

REPERENCE/DOCKET NUMBER: 33,730

TELLEPAX: (415) 327-3231

INFORMATION POR SEG DID NO: 10: SEQUENCE CHARACTERISTICS:

LENGTH: 943 amino acids

TURER ADDRESSED ANDREMED ACIDS

TOCKET TO ACIDS

TOCKET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 48.4%; Pred. No. 0.52;
Matches 15; Conservative 5; Mismatches 8; Ind
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; TOPOLOGY: linear; MOLECULE TYPE: protein US-08-455-970A-10
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                                                                                                                             Application US/08676263
                                            98 AGCKNFYWKGFTSC 111
                                                         1 AGCKNFFWKTFTSC
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US-08-676-263-11 ; Sequence 11, A
Query Match
Best Local S
Matches 12
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COMPOSITIONS AND METHODS FOR THE DELIVE OF DRUGS BY PLATELETS FOR THE TREATMENT CARDIOVASCULAR AND OTHER DISEASES
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                                                                                                                                                                                                                                                                             TYPE: Floppy disk

TYPE: Floppy disk

FER: IBM PC compatible

FING SYSTEM: PC-DOS/MS-DOS

ARE: PatentIn Release #1.0, Version

APPLICATION DATA:
                                                                                                                                                                 O'Toole, Gerstein,
ker Drive/6300 Sea
                                                                                                                                                                                                                                                                                                                                                                     MBER: US/08/676,263
07-NOV-1996
                                                                                                                                                                                                                                  United States of America 6-6402
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2-JAN-1994
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Franklin Street
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ADDRESSEE: Fish & Ric
STREET: 225 Franklin
CITY: Boston
STATE: Massachusetts
                                                                                                                                                                                                                                                                    FORM:
                                                                                                                                                                 Marshall
                                                                                                                                                                                                                                                                                                                                                                       CATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER:
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NUMBER OF SEQUENCES
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12; Conser
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APPLICANT: Vint
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US-08-676-263-11
                                                                                                                                                                                                                                   COUNTRY:
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Best Local S
Matches 12
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US-08-286-
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                                                                                                                                                               Length
                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Vincent, Jean-Pierre
APPLICANT: Gaudriault, Georges
APPLICANT: Beaudet, Alain
TITLE OF INVENTION: FLUORESCENT SOMATOSTATIN
NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
                                                                                                                                                              Score 77; DB 1
Pred. No. 0.002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 06942/003001
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
TELEFAX: 617/542-8906
TELEX: 200154
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 14 amino acids
TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 77; DB Pred. No. 0.001; Mismatches
                                                                                                                                                                                          Mismatches
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APPLICATION NUMBER: US/08/416,007 FILING DATE: 04-APR-1995 CLASSIFICATION: 530 ATTORNEY/AGENT INFORMATION: NAME: Clark, Paul T. REGISTRATION NUMBER: 30,162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: USA
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, V
                                                                                                                                                                                                                                                                                                                                                 Sequence 4, Application US/08416007
Patent No. 5693679
GENERAL INFORMATION:
APPLICANT: Vincent, Jean-Pierre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      not relevant
                                                                                                                                                         th 12.9%; similarity 85.7%; 12; Conservative
; SEQUENCE CHARACTERISTICS:
; LENGTH: 14 amino acids
; TYPE: amino acid
; TOPOLOGY: unknown
; MOLECULE TYPE: peptide
; HYPOTHETICAL: NO
; ANTI-SENSE: NO
US-08-255-272-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       th 12.9%; Similarity 85.7%; 12; Conservative
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TOPOLOGY: linear

MOLECULE TYPE: protein

US-08-416-007-4
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Boston

STATE:

RESULT 13 US-08-416-007-4

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Query Match Best Local S Matches 12

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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
COMPUTER: IBM PS/2 Model 502 or 55SX
OPERATING SYSTEM: MS-DOS (Version 5.0)
SOFTWARE: WordPerfect (Version 5.1)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/286,748B
FILING DATE: August 5, 1994
CLASSIFICATION DATA:
APPLICATION NUMBER: 32,983
FILING DATE: ATTORNEY/AGENT INFORMATION:
NAME: J. Peter Fasse
REGISTRATION NUMBER: 32,983
REGISTRATION NUMBER: 32,983
REGISTRATION NUMBER: 32,983
REFERENCE/DOCKET NUMBER: 04547/013001
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEX: 200154
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 14
TYPE: amino-acid
STRANDEDNESS: single
TOPOLOGY: linear
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Search completed: March 21, 2003, 11:42:29. Job time: 21.9614 secs ОЪ

Length

Score 77; DB 1; Pred. No. 0.002; 1; Mismatches

Ouery Match 12.9%; Best Local Similarity 85.7%; Matches 12; Conservative

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Ogjrgg neisserla m
Ogldc2 oryza sativ
Ogld76 homo sapien
O75175 homo sapien
Oglrf2 oryza sativ
O54501 streptomyce
Oguhk3 homo sapien
Ogkz43 streptomyce
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Q9epq3 mus musculu
Q9hn51 halobacteri
Q9sbx3 zea mays (m
Q9fer7 zea mays (m
                                                                                                                                                                                                                                                                                                                                   OO7834 pseudomonas
O9m6d8 oryza sativ
O9p4a6 letharia vu
O8zph5 salmonella
O9agi4 pseudomonas
O9hvr9 pseudomonas
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Ostariophysi; Cypriniformes
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Al-Mahrouki A.A., Irwin D.M., Youson J.H.;
"Molecular cloning and characterization of white sucker preprosomatostatin.";
Submitted (AUG-2000) to the EMBL/GenBank/DDBJ databases EMBL; AF292654; AAK97071.1;
InterPro, IPR004250; Somatostatin.
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00828D35263E8805 CRC64;
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Last annotation update)
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Pred. No. 7.1e-08;
5; Mismatches 18
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WTKKDIEELLSQLSLPEIEAREN 75
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Chordata, Craniata;
oterygii; Teleostei;
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A; 13783 MW;
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Catostomidae; Catostomus.
NCBI_TaxID=7971;
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                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 114;
                                                                                                                                                                                            Al-Mahrouki A.A., Irwin D.M., Youson J.H.; "Characterization of variant somatostatin cDNAs from several osteoglossomorphs: molecular identification and comparative analysis.":
                                                                                          Osteoglossomorpha;
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--RQLLQRTLAAS
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eleostei; Osteoglosso
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aryota; Metazoa; Chordata; Craniata; Vertebi
inopterygil; Chondrostel; Acipenseriformes;
                                                                                                                                                                                                                                                                                                                                                                                                                             Score 93.5; DB 1 Pred. No. 0.0045;
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                                                                                                                                                                                                                                                                                         Submitted (AUG-2000) to the EMBL/Ge
EMBL; AF292652; AAK97069.1; -.
InterPro; IPR004250; Somatostatin.
Pfam; PF03002; Somatostatin; 1.
SEQUENCE 114 AA; 12494 MW; 4541
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S; Mormyridae;
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116 AA; 12616 MW:
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NCBI_TaxID=42645;
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Trabucchi M.,
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1-MAR-2002
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090Y40;
01-DEC-2001
01-JUN-2002
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O90Y43
O90Y43;
O90Y43;
O1.DEC-2001 (TrEMBLrel. 19, Created)
O1.DEC-2001 (TrEMBLrel. 19, Last sequence update)
O1.DEC-2001 (TrEMBLrel. 20, Last annotation update)
O1.DEC-2001 (TrEMBLrel. 20, Last annotation update)
O1.MAR-2002 (TrEMBLrel. 20, Last annotation update)
O1.MACITY (Tremble Constant)
O1.MAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
Al-Mahrouki A.A., Irwin D.M., Youson J.H.;
Al-Mahrouki A.A., Irwin D.M., Youson J.H.;
"Characterization of variant somatostatin cDNAs from several costeoglossomorphs: molecular identification and comparative analysis.";
Submitted (AUG-2000) to the EMBL/GenBank/DDBJ databases.
EMBL; AF292651; AAK97068.1;
InterPro; IPR004250; Somatostatin.
Pfam; PF03002; Somatostatin; 1.
SEQUENCE 114 AA; 12352 MW; 7E3D44CB6A27B12F CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       090v42
090v42
090v42;
01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-DEC-2002 (TrEMBLrel. 20, Last annotation update)
01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
Preprosomatostatin.
Pantodon buchholtzi (Butterflyfish).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutela Actinopterygii; Neopterygii; Teleostei; Osteoglossomorpha Osteoglossiformes; Pantodontidae; Pantodon.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
24.9%; Score 112; DB 13;
Best Local Similarity 56.5%; Pred. No. 3.6e-05;
Matches 26; Conservative 4; Mismatches 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
24.0%; Score 108; DB.13;
Best Local Similarity 50.0%; Pred. No. 0.0001;
Matches 23; Conservative 9; Mismatches 12;
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RESULT Q90Y42

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RESULT 4 090Y41 ID 090Y AC 090Y DT 01-D DT 01-D

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MEDLINE-21082930;
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PF00535;
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Best Local S
Matches 29
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  114;
                                                                                                                                                                                                                   several
                                    Osteoglossomorpha;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 50.0%; Pred. No. 0.13;
Matches 22; Conservative 6; Mismatches 13; Indels
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OBXXAO;

O1-MAR-2002 (TrEMBLrel. 20, Last sequence update)

O1-MAR-2002 (TrEMBLrel. 21, Last annotation update)

O1-JUN-2002 (TrEMBLrel. 21, Last annotation update)

O1-JUN-2002 (TrEMBLrel. 21, Last annotation update)

DNAG OR RSC2216 OR RS01383.

Ralstonia solanacearum (Pseudomonas solanacearum).

Bacteria; Proteobacteria; beta subdivision; Ralstonia grr Ralstonia.

NCBL_TAXID=305;

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SEQUENCE FROM N.A.

SEQUENCE FROM N.A.

SEQUENCE FROM N.A.

SEQUENCE FROM N.A.

SALAIN-EMILOUD;

A Arlat M., Billault A., Brottier P., Camus J.C., Cattolic Achandler M., Choisne N., Claudel-Renard C., Cunnac S., DA Arlat M., Choisne N., Claudel-Renard C., Saurin W., Sci Arlat M., Walsan M., Molsan A., Robert C., Saurin W., Sci Aslanoubat M., Molsan A., Robert C., Saurin W., Sci Siguier P., Thebault P., Whalen M., Wincker P., Levy M., Weissenbach J., Boucher C.A.;

Neissenbach J., Boucher C.A.;

Siguier P., Thebault P., Whalen M., Wincker P., Levy M., Mature 415:497-502(2002)

EMBL; AL646068; CAD15923.1;

InterPro; IPR002936; DNAprim, 1.

REMBL; RM0493; TOPRIM; 1.

REMBL; SM00400; Zf-CHC2; 1.

SMART; SM00400; Zf-CHC2; 1.

SMART; SM00400; Zf-CHC2; 1.

SEQUENCE 603 AA; 66305.MW; 5A3A29A1674E541C CRC64;
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                                                                                                                                          SEQUENCE FROM N.A.
Al-Mahrouki A.A., Irwin D.M., Youson J.H.;
Al-Mahrouki A.A., Irwin D.M., Youson J.H.;
"Characterization of variant somatostatin cDNAs from osteoglossomorphs: molecular identification and compa analysis.";
Submitted (AUG-2000) to the EMBL/GenBank/DDBJ databas EMBL; AF292653; AAK97070.1; -.
InterPro; IPR004250; Somatostatin.
Pfam; PF03002; Somatostatin; 1.
SEQUENCE 114 AA; 12561 MW; 4E3C32F58E34F971 CRC64
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Pred. No. 1.2;
3; Mismatches 15; Ind
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Actinopterygii; Neopterygii; Teleostel; O:
Osteoglossiformes; Notopteridae; Chitala.
NCBI_TaxID=112163;
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Best Local Similarity 38.6%;
Matches 22; Conservative
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ID QBXXA

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DT O1-1

DT O1-1

AC BAC

OC RA

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VSQIHCALALLGLALAICSQGAASQPDLDLASRRLLQRALAAALPHRSGVSERWRTFYPN
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                                                        Rhizobium loti (Mesorhizobium loti).
Bacteria; Proteobacteria; alpha subdivision; Rhizobiaceae
Phyllobacteriaceae; Mesorhizobium.
NCBI_TaxID=381;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  subdivision; Xanthomonas
                                                                                                                                                                                                                                                                                                                  01173; Glycos_transf_2.
Glycos_transf_2; 1.
cotein; Complete proteome.
AA; 27987 MW; 45709C6A71EDD2AA CRC64;
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Nakazaki N., Shimpo
                                                                                                                                                                                                                                                                                                                                                                                              16;
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Last annotation up
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Mismatches
Last sequence up
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Pred. No.
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AP003000; BAB49897.1;
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29; Conservative
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Bacteria; Proteobacteria;
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251 AA; 2
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Ouery Match
Best Local S
Matches 18
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093W36;
01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 20, Last annotation update)
01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
At2904160/T16B23.1.
Arabidopsis thaliana (Mouse-ear cress).
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnollophyta; eudicotyledons; core eudicots; Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
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Shinn P., Chenk R., Kim C.J., Koesema E., Meyers M.C.,
Banh J., Bowser L., Carninci P., Dale J.M., Gibson H.A.,
Banh J., Bowser L., Carninci P., Dale J.M., Gibson H.A.,
Goldsmith A.D., Hayashizaki Y., Ishida J., Jiang P.X., Jones T.,
Kamiya A., Kariin-Neumann G., Kawai J., Lam B., Lee J.M., Lin J.,
Liu S.X., Miranda M., Narusaka M., Nguyen M., Onodera C.S., Palm C.
Pham P.K., Quach H.L., Sakurai T., Satou M., Seki M., Southwick A.,
Tang C.C., Toriumi M., Yamada K., Yu G., Yu S., Shinozaki K.,
A Davis R.W., Theologis A., Ecker J.R.;
Submitted (JUL-2001) to the EMBL/GenBank/DDBJ databases.
EMBL; AYOS8235; AAL15409.1; -
EMBL; AYOS8235; AAL15409.1; -
EMBL; AYOS8235; PAL16405.1; -
InterPro; IPRO01377; PA.
InterPro; IPRO0209; Peptidase_S8.
Fram; PF00082; Peptidase_S8.
Pfam; PF00082; Peptidase_S8; 1.
RPfam; PF00082; Peptidase_S8; 1.
RPfam; PF00082; Peptidase_S8; 1.
RPGAM; PF00082; Peptidase_S8; 1.
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Narusaka
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Nhani A. Jr., Nobrega F.G., Nunes L.R., Oliveira M.A.,

de Oliveira M.C., de Oliveira R.C., Palmieri D.A., Paris A.,

Peixoto B.R., Pereira G.A.G., Pereira H.A. Jr., Pesquero J.B.,

Quaggio R.B., Roberto P.G., Rodrigues V., de Rosa A.J.M.,

de Rosa V.E. Jr., de Sa R.G., Santelli R.V., Sawasaki H.E.,

da Silveira J.F., Silvestri M.L.Z., Siqueira W.J., de Souza A.A.

de Souza A.P., Terenzi M.F., Truffi D., Tsai S.M., Tsuhako M.H.,

Vallada H., Van Sluys M.A., Verjovski-Almeida S., Vettore A.L.,

Zago M.A., Zatz M., Meidanis J., Setubal J.C.;

"The genome sequence of the plant pathogen xylella fastidiosa."

R PREL, AE04055; AAF85270.1;

R InterPro; IPR002604; ATZ_TRZ.

R Pfam; PF01685; ATZ_TRZ.

R Pfam; PF01685; ATZ_TRZ.

Hypothetical protein; Complete proteome.

SEQUENCE 435 AA; 47299 MW; E9272E57460CB447 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kim C.J., Chen H., Cheuk R., Koesema E., Meyers M.C., Banh J., Kim C.J., Chen H., Cheuk R., Koesema E., Meyers M.C., Banh J., Bowser L., Carninci P., Dale J.M., Goldsmith A.D., Hayashizaki Ishida J., Jiang P.X., Jones T., Kamiya A., Karlin-Neumann G., Kawai J., Lam B., Lee J.M., Lin J., Liu S.X., Miranda M., Narus Nguyen M., Onodera C.S., Palm C.J., Pham P.K., Quach H.L., Saku Satou M., Seki M., Southwick A., Tang C.C., Toriumi M., Yamada Yamamura Y., Yu G., Yu S., Shinozaki K., Davis R.W., Theologis Ecker J.R.; "Arabidopsis ORF clones."; "Arabidopsis ORF clones."; Submitted (OCT-2001) to the EMBL/GenBank/DDBJ databases.
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                                                                                                                                                                                                                                                                                                                                                                                                                           15;
                                                                                                                                                                                                                                                                                                                                                                                   435;
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Best Local Similarity 33.9%; Pred. No. 12;
Matches 20; Conservative 6; Mismatches
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Q93W36
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Embryophyta; Tracheophyta; dons; core eudicots; Rosidae;
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae;
eurosids, II; Brassicales; Brassicaceae; Arabidopsis.
                                                   -ALPHRSGVSERWRTFYPNCPCLRWRP
                                                                             -WSP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
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J.C.;
to the EMBL/GenBank/DDBJ databases
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L., Ronning C.M.
, Town C.D., Nie
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Spermatophyta; Magnoliophyta; eudicotyledons; core
eurosids II; Brassicales; Brassicaceae; Arabidopsis
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ase AIR3 (Fragment)
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18;
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Pred. No. 18;
13; Mismatches
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T.M., Bowman C.L.
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Score 68;
Pred. No.
                                                   GLALAICSQGAASQPDLDLASRRLLQRALAA
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                                                                                                                                                                                                                         01-JUN-2002 (TrEMBLrel. 21, Crea 01-JUN-2002 (TrEMBLrel. 21, Last 01-JUN-2002 (TrEMBLrel. 21, Last Subtilisin-like serine protease AT2G04160.
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                         13;
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28.18;
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AC007178; AAM15440.1
                       Conservative
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Best Local Similarity 28.1
Matches 18; Conservative
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08S896;
01-JUN-2002 (TrEMBLrel.
01-JUN-2002 (TrEMBLrel.
01-JUN-2002 (TrEMBLrel.
Subtilisin-like serine parzg04160.
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Barnstead M.E., Mason
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                                                                                                                                                                                                                                                                                                                                                                                    COLUMBIA;
          Local Similarity
les 18; Conser
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Kaul
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Eukaryota; V
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STRAIN=CV.
Lin X., Ka
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Barnstead
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Q8S8B0;
01-,777
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; Tracheophyta; eudicots; Rosidae;

update) on update)

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Arabidopsis thaliana (Mouse-ear cress).
Eukaryota; Viridiplantae; Streptophyta; Embryophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eleurosids II; Brassicales; Brassicaceae; Arabidopsis.
NCBI_TaxID=3702;
                                          -MAY-1999 (TrEMBLrel. 10, Last sequence up-JUN-2002 (TrEMBLrel. 21, Last annotation
                                                                                                    Subtilisin-like protease AIR3.
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TO 1-NOV-1998 (TTEMBLrel. 08, Created)

TO 1-NOV-1998 (TTEMBLrel. 08, Last sequence update)

TO 1-NOV-1998 (TTEMBLrel. 12) Last annotation update)

DT 01-UND-2002 (TTEMBLrel. 21, Last annotation update)

E Subtlisin-like protesse (Fragment).

SUBTLISIN-like protesse (Fragment).

SUBTLISIN-like protesse (Fragment).

SECONDERS (SECONDERS)

C EUKARYOTA: Viridiplantae (Mouse-ear Cress).

C EUKARYOTA: Viridiplantae (Mouse-ear Cress).

C EUKARYOTA: Viridiplantae (Mouse-ear Cress).

C EUKARYOTA: Viridiplantae: Streptophyta; Embryophyta; Tracheophyta;

C EUKARYOTA: Viridiplantae: Streptophyta; Embryophyta; Tracheophyta;

C EUKARYOTA: Viridiplantae: Streptophyta; Embryophyta; Tracheophyta;

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Best Local Similarity 28.1%; Pred. No. 24;
Matches 18; Conservative 13; Mismatches 25; Indels
Fraser C.M., Venter J.C.;
Submitted (FEB-2002) to the EMBL/GenBank/DDBJ databases
EMBL; AC007293; AAM15483.1; •.
Protease.
NON_TER 755 755
SEQUENCE 755 AA; 80844 MW; 2E9BE365795936DE CRC64;
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Best Local Similarity 28.1%; Pred. No. 24;
Matches 18; Conservative 13; Mismatches 25; Ind
                                                                                                                                                                                                                                                                                                                                                           Q9ZSP5 PRELIMINARY; PRT; Q9ZSP5; 01-MAY-1999 (TrEMBLrel. 10, Created)
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584 AAIR 587
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Q9ZSP5
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T., Nhan M.
                                                         Hooykaas
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                                                                               Arabidopsis
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AE008824; AAL21627.1; -.
Metical protein; Complete proteome.
MCE 197 AA; 22326-MW; 7CA059444F962F66 CRC64;
                                                                                                                                                                                                                                                         75DCCA0DED63F47E CRC64;
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01-MAR-2002 (TrEMBLrel. 20, Last sequence update)
01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
Putative periplasmic protein.
                                                        O.R.,
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Stoneking
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SEQUENCE FROM N.A.
STRAIN-CV. COLUMBIA;
MEDLINE-99246056; PubMed-10231025;
MEDLINE-10246056; PubMed-10231025;
MEDLINE-99246056; PubMed-10231025;
MEDLINE-99246056; PubMed-10231025;
MEDLINE-99246056; PubMed-10231025;
                                                                                  sase gene from A root emergence.
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InterPro; IPRUCCE.

Pfam; PF02225; PA; 1.

Pfam; PF00082; Peptidase_S8; 2.

Pfam; PF000723; SUBTILISIN.

PRINTS; PR00723; SUBTILISIN.

PRINTS; PR00138; SUBTILASE_SER; UNKNOWN_1
                                                                                                                                                                                                                                                                                                                  Mismatches
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Sanderson K.E., Spier
Sanderson K.E., Spier
Sanderson K.E., Holmes
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Pred. No.:
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Pred.
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Miller
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                                                                                "A novel subtilisin-like protease expressed at sites of lateral root DNA Res. 6:13-19(1999).
EMBL; AF098632; AAD12260.1; -.
                                                                                                                                                      IPR003137; PA.
IPR000209; Peptidase_S8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       gamma
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H., Florea L., N
Wilson R.K.;
Nome sequence of
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Bacteria; Proteobacteria;
Salmonella.
                                                                                                                                                                                                                                                                                                                  Conservative
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STRAIN-LT2 / SGSC1412
MEDLINE-21534948; Pubh
                                                                                                                                                                                                                                                                                          1 Similarity
18; Conserve
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"Complete genome
LT2 ".
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EMBL; AE00882
Hypothetical
SEQUENCE 19
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08ZMR3;
01-MAR-2002
01-MAR-2002
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SEQUENCE
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Matches 18
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Search completed: March 21, 2003, 11:40:43 Job time : 46.0773 secs

us-09-727-739b-17

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Title:
US-09-727-739B-17
Perfect score: 450
Sequence: 1 MRVSQIHCALALLGLALAIC.....RWRPRKVKGPQLKAKEDLER 86
                                                                                                         March 21, 2003, 11:35:23; Search time 10.3348 Seconds (without alignments) 345.142 Million cell updates
GenCore version 5.1.4_p5_4578 Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                    OM protein - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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112892 Total number of hits satisfying chosen parameters: 112892 seqs, 41476328 residues . Searched:

Minimum DB seq length: 0 Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

SwissProt_40:* Database :

Pred. No. is the number of results predicted by chan score greater than or equal to the score of the resu and is derived by analysis of the total score distri

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rhodothermu	homos	barley mile	mus mu	caenorhabdi	human cytom	saccharomyc	homo sapien	homo s	homo s	azotobacter	homo s
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DNAB_RHOMR	TRL2_HUMAN	POL2_BAMMU	E4L2_MOUSE	E2K3_CAEEL	UL50_HCMVA	ORC5_YEAST	MSH5_HUMAN	SPA1_HUMAN	SPCR_HUMAN	NIFQ_AZOVI	DLX4_HUMAN
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941	1503	894	. 886	1077	397	479	834	1042	3674	195	347
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28	58	57.5	57.5	57.5	57	57	57	57	57	56.5	56.5
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RESULT SMS2 O	JLT 1 2 ONCMY
ID.	SMS2_ONCMY STANDARD; PRT; 115 AA.
Z E	Q91194; 16-00m-2001 (Do) 40 Created)
	16-OCT-2001 (Rel. 40, Last sequence update)
	tatio
	ains:
	Oncorhynchus mykiss (Rainbow trout) (Salmo gairdneri).
88	raniata; Vertebrata;
	mos. calmontan.
3 8	
RN	
RP	QUENCE FROM N.A.
RX	54921; PubMed-7628684;
RA	Kittilson J.D., Dahl S.K., Sheridan M.
RT	and characterization of a cDNA encoding for
RI	preprosomatostatin containing [Tyr7, Gly10]-somatostatin-14 from the
T I	valicities of failings tiout, outdothylichus mykiss podooring 00.051-061/1005)
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3 5	: SUBCELLULAR LOCATION: Secreted.
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ပ္ပ	SWISS-PROT entry is copyright. It is produced through
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ပ္ပ	European Bioinformatics Institute. There are no restri
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2	an email to license@isb-sib ch)
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DR	F03002; Somatostatin; 1.
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OS:	QUENCE 115 AA; 12963 MW;
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œ ,	Query Match No. 2.2e-32: Dest. No. 2.2e-32: Dest. Local Similarity 85.7%; Pred. No. 2.2e-32:
X	72; Conservative 2; Mis
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ò	61 DNCPCI.RWRDRKVKGPOI.KAKEDI. 84
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39b-17

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hormone; Signal; Multigene family
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Euteleostomi;
dae; Phasianinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         There are no restrictions or as its content is in read of the content is in read of the content is in 
   Euteleostom1;
; Cypriniformes;
                                                                                                                                                                                                encoding preprosomatostatin-I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-OCT-1993 (Rel. 27, Created)
01-OCT-1993 (Rel. 27, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Somatostatin precursor [Contains: Somatostatin-14]
                                                                                                                                                                                                                                                                                                                                                           is produced through a collabrantics and the EMBL out:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sib.ch
                                                                                                                                                                                 "Cloning and characterization of cDNAs encoding preprosomatostat and -II from goldfish brain.";
Submitted (NOV-1995) to the EMBL/GenBank/DDBJ databases.
-!- FUNCTION: SOMATOSTATIN INHIBITS THE RELEASE OF SOMATOTROPIN.
-!- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   to the EMBL/GenBank/DDBJ databases.
TATIN INHIBITS THE RELEASE OF SOMATOTROPIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 114;
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                                                                                                                                                                                                                                                                                                                                                                                                            There are no rest
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Galliformes; Phasianidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Karahashi K., Kato S., Yamamoto H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOMATOSTATIN-26 (POTENTIAL)
SOMATOSTATIN-14.
BY SIMILARITY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CRC64;
Craniata; Vertebrata; El
Teleostel; Ostariophysi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Usage by http://www
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  http://www
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TISSUE-Pancreas;
Nata K., Kobayashi T., Karahashi K., Kato S., Yamam Yonekura H., Okamoto H.;
Submitted (JUN-1991) to the EMBL/GenBank/DDBJ datak-1- FUNCTION: SOMATOSTATIN INHIBITS THE RELEASE OF-1- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        42
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                                                                                                                                                                                                                                                                                                                                                        This SWISS-PROT entry is copyright. It is produbetween the Swiss Institute of Bioinformatics the European Bioinformatics Institute. There a use by non-profit institutions as long as impositied and this statement is not removed. Use entities requires a license agreement (See http. or send an email to license@lsb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 91.5; DB 1
Pred. No. 0.0016;
5; Mismatches 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  04250; Some Somatostatin; 1.
Somatostatin; 1.
ir of basic residues; Horm POTENTIAL.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRT;
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Eukaryota; Metazoa; Chordata; (
Archosauria; Aves; Neognathae;
Eukaryota; Metazoa; Chordata;
Actinopterygii; Neopterygii; '
Cyprinidae; Carassius.
NCBI_TaxID=7957;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12574 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20.3%;
ilarity 54.8%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            54; AAD09359.1;
IPR004250; Somat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STANDARD;
                                                                                                                                                                         R.E.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
                                                                                                                   SEQUENCE FROM N.A TISSUE-Brain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              usuU2; Sc
on pair
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25
89
101
103
114 A
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                                                                                                                                                                      Lin X.-W.,
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DISULFID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EMBL;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      omatostatin-I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           estrictions
                                                                                                                                                                                                                                                                                       teleostomi;
Cypriniforme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TISSUE-Liver;
Otto C.J., Peter R.E.;
The expression of SRIF mRNA in the brain of goldfish.";
Submitted (SEP-1997) to the EMBL/GenBank/DDBJ databases.
-!- FUNCTION: SOMATOSTATIN INHIBITS THE RELEASE OF SOMATOTROPIN.
-!- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY.

-!- SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOTROPIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   residues; Hormone; Signal; Multigene POTENTIAL.
POTENTIAL.
[TYR21,GLY24]SOMATOSTATIN-28.
[TYR7,GLY10]SOMATOSTATIN-14.
BY SIMILARITY.
O -> RW (IN REF. 2).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     120;
                                                                                    SMS2_CARAU STANDARD; PRT; 120 AA.

O9YGH4; Q9PTU2;
16-OCT-2001 (Rel. 40, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
16-OCT-2001 (Rel. 40, Last annotation update)
16-OCT-2001 (Rel. 40, Last annotation update)
Somatostatin II precursor [Contains: [Tyr21,Gly24]somatost
[Tyr7,Gly10]somatostatin-14].
Carassius auratus (Goldfish).
Eukaryota; Metazoa; Chordata; Cranlata; Vertebrata; Eutele
Actinopteryg11; Neopteryg11; Teleoste1; Ostarlophys1; Cypr
NCBI_TaxID=7957;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Lin X.-W., Peter R.E.; "Cloning and characterization of cDNAs encoding prepros and -II from goldfish brain."; Submitted (JUN-1996) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                )6;
11; Indel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  -> RW (IN REF. 2).
98957D68011A651A CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            update)
Somatostatin-26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 MRVSQIHCALALLGLALAICSQGAASQ--PDLDLASRRLLQRALA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 red. No. 1.3e-06;
Mismatches 11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         114 AA
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16-OCT-2001 (Rel. 40, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Somatostatin IA precursor [Contains: Somate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 119;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EMBL; U60262; AAD09626.1; -
EMBL; AF025686; AAF15306.1; -
InterPro; IPR004250; Somatostatin.
Pfam; PF03002; Somatostatin; 1.
Cleavage on pair of basic residues; 1
SIGNAL 1 23 POTENTI
PROPEP 24 92 POTENTI
PROPEP 93 120 [TYR21, 0]
PEPTIDE 107 120 BY SIMII
CONFLICT 51 51 0 -> RW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              13723 MW;
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Carassius auratus (Goldfish)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26.4%;
ilarity 57.8%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24
93
107
109
51
120 AA;
                                                                                                                                                                                                                                                                                                                                                                                                           FROM N.A.
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26; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                    TISSUE-Brain;
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09YGH5;
16-OCT-2001
16-OCT-2001
16-OCT-2001
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PEPTIDE
DISULFID
CONFLICT
SEQUENCE
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                                     RESULT 2
SMS2_CARAU
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Matches
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Siluriformes
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                                                                                                                                                                                                                                                                                                                            pancreatic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Dixon J.
                                                                                                                                                                                                                                                                                                .13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SMS1_ICTPU STANDARD; PRT; 114 AA.

SMS1_ICTPU STANDARD; PRT; 114 AA.

21-JUL-1986 (Rel. 01, Created)

01-AUG-1990 (Rel. 15, Last sequence update)

16-OCT-2001 (Rel. 40, Last annotation update)

Somatostatin I precursor [Contains: Somatostatin-14 (SS-1)

Ictalurus punctatus.(Channel catfish).

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutel

Actinopterygii; Neopterygii; Teleostei; Ostariophysi; Sill

Ictaluridae; Ictalurus.

NCBI_TaxID=7998;
                                                                                                                                                                                                                                                                                              Indels
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or send an email

EMBL; X60191; CAA42747.1; -.

EMBL; X60191; CAA42747.1; -.

PIR; S20630; S20630.

InterPro; IPR004250; Somatostatin.

Reavage on pair of basic residues; Hormone; Signal.

T Cleavage on pair of basic residues; Hormone; Signal.

T PROPEP 25 88 BY SIMILARITY.

T PROPEP 25 88 BY SIMILARITY.

T PEPTIDE 103 116 SOMATOSTATIN-14.

TT PEPTIDE 103 116 SOMATOSTATIN-14.

THE T AA; 12675 MW; BA5BB9BDA8A291BA CRC64

116 105 116 105 116

117 AA; 12675 MW; BA5BB9BDA8A291BA CRC64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE-82265698; PubMed=6179939;
Minth C.D., Taylor W.L., Magazin M.D., Tavianini M.A.,
Weith H.L., Dixon J.E.;
"The structure of cloned DNA complementary to catfish
somatostatin-14 messenger RNA.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE-81264223; PubMed-6114953;
Andrews P.C., Dixon J.E.;
"Isolation and structure of a peptide hormone predicte sequence. A second somatostatin from the catfish panci J. Biol. Chem. 256:8267-8270(1981).
-!- FUNCTION: SOMATOSTATIN INHIBITS THE RELEASE OF SOM-1- SUBCELLULAR LOCATION: Secreted.
-!- TISSUE SPECIFICITY: PANCREAS.
-!- SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY.
                                                                                                                                                                                                                             8A5BB9BDA8A291BA CRC64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE OF 82-114 FROM N.A. MEDLINE-82082515; PubMed-6171821; Taylor W.L., Collier K.J., Deschenes R.J., Weith H.L. "Sequence analysis of a cDNA coding for a pancreatic
                                                                                                                                                                                                                                                     Score 83.5; DB 1;
Pred. No. 0.013;
14; Mismatches 25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 somatostatin.";
Proc. Natl. Acad. Sci. U.S.A. 78:6694-6698(1981)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.
MEDLINE=85303576; PubMed=2863931;
Dixon J.E., Andrews P.C.;
"Somatostatins of the channel catfish."
Adv. Exp. Med. Biol. 188:19-29(1985).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  messenger RNA."; 257:10372-10377(1982).
                                                                                                                                                                                                                                                                                                                                                                                            64 PCLRWRPRKVKGPQLKAKEDLER 86
| | : |: || |
56 --LLSEPSQTENEALES-EDLSR 75
                                                                                                                                                                                                                                                   18.6%;
al Similarity 37.3%;
31; Conservative 14
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Matches 31
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collaboration
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                                                            oved. Usage by and for commercial (See http://www.isb-sib.ch/announce/
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                           restrictions on
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s octreotide or SMS
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ne treatment of a variety of disorders
// and the symptomatic treatment of carcinoid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (Cynomolgus monkey).
rtebrata; Euteleostomi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ion-enhanced DNA-driven subtractive cDNA cloning: low-abundance monkey cortex-specific mRNAs. "; cad. Sci. U.S.A. 85:1696-1700(1988). SOMATOSTATIN INHIBITS THE RELEASE OF SOMATOTROPIN.
                                                          for
                                                                                                                                                                                                                                                                                                                                       Length 114;
through
                                                                                                                                                                                                                                                                                                                                                                    Indels
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FEE0F2C76F74D99F CRC64;
          Bioinformatics and titute. There are no
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is produced
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tion update)
Somatostatin-28;
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Shen L.-P., Pictet R.L., Rutter W.J.;
"Human somatostatin I: sequence of the cDNA.";
Proc. Natl. Acad. Sci. U.S.A. 79:4575-4579(1982)
                                                                                                                                                                                                                                                                                                                                     Score 80.5; DB 1;
Pred. No. 0.028;
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SOMATOSTATIN-14
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                                 long
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Chordata; Craniata; Vo
Primates; Catarrhini;
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                                         use by non-profit institutions as lo modified and this statement is not remoentities requires a license agreement (or send an email to license@lsb-slb.ch)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1-JUL-1986 (Rel. 01, Created)
1-JUL-1986 (Rel. 01, Last sequence up
6-OCT-2001 (Rel. 40, Last annotation
somatostatin precursor (Contains: Some
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                          Institute
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IEDLINE=88144503; PubMed=2894033;
ravis G.H., Sutcliffe J.G.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PubMed=6142531;
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V00607; CAA23877.1;
V00608; CAA23878.1;
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Eukaryota; Metazoa; Che
Mammalia; Eutheria; Pr
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21; Conser
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MEDLINE-84146798;
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P01166;
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SMS_HUMAN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOMATOTROPIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               family;
                                                                                                                                                                                                               Multigene family;
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                                                                                                                                                                                                                                                                                                                Length 115;
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(PSS1) [Contains: Somatostatin-27
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Lepidosireniformes; Protopteridae; Protopterus
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Bioinformatics and
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SOMATOSTATIN INHIBITS THE RELEASE OF
                                                ht. It is produced
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S
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SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY
                                                                                                                        http:/
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349756FEB4ABE213
                                                                                                                                                                                                                                                                                                                 1;
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r of basic residues; Hormone, 24 BY SIMILARITY.
99 BY SIMILARITY.
99 SOMATOSTATIN-14.
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SOMATOSTATIN-14
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Tostivint H., Lihrmann I.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 lungfish)
Craniata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the European and Institutions as modified and this statement is not remoentities requires a license agreement (entities requires to licenselish-sib.ch)
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                                                                                                                        a license agreement (to license@lisb-sib.ch)
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Last annotation
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Pred.
                                               This SWISS-PROT entry is copyright. between the Swiss Institute of Bicthe European Bioinformatics Institutuse by non-profit institutions amodified and this statement is not entities requires a license agreement.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              entry is copyright
as Institute of Bio
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12691 MW;
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the European Bioinformatics
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(Rel. 40,
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22; Conserv
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MEDLINE-99326690;
Trabucchi M., Tost
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Pfam; PF03002; Sc
Cleavage on pair
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104
115 A
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NCBI_TaxID=7888
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Pfam; PF03002;
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P87384; Q9PS18;
16-OCT-2001 (Rel. 40, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
16-OCT-2001 (Rel. 40, Last annotation update)
Somatostatin 1 precursor (PSS1) [Contains: Somatostatin-14 (S-1) (SSS1)].
Rana ridibunda (Laughing frog) (Marsh frog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Amphibia; Batrachia; Anura; Neobatrachia; Ranoidea; Ranidae; Rane
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family
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SEQUENCE FROM N.A.
TISSUE-Brain;
MEDLINE-97057290; PubMed-8901629;
Tostivint H., Lihrmann I., Bucharles C., Vieau D., Coulouarn Fournier A., Conlon J.M., Vaudry H.;
"Occurrence of two somatostatin variants in the frog brain: characterization of the cDNAs, distribution of the mRNAs, and receptor-binding affinities of the peptides.";
Proc. Natl. Acad. Sci. U.S.A. 93:12605-12610(1996).
                                                                                                                                                                                                                                                                                                                                                                                                        116;
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Vaudry H., Chartrel N., Conlon J.M.;
"Isolation of [Pro2,Met13]somatostatin-14 and somatostat
frog brain reveals the existence of a somatostatin gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        chem. Biophys. Res. Commun. 188:477-482(1992).
FUNCTION: SOMATOSTATIN INHIBITS THE RELEASE OF SOMAT
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                                                                                                                                                                                                                                                                                                                                                                                                       Length
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Pred. No. 0.037;
13; Mismatches 28;
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| | | : : | : | | | 56 --LLSEPNQTENDALE-PEDLSQ 75
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Conservative 1
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TJ-DEC-1998 (Rel. 37, Last sequence update)

TJ-DEC-1998 (Rel. 37, Last annotation update)

SGM SOMALOSTAIN PECURSOR [Contains: Somatostatin-28; Somatostatin precursor [Contains: Somatostatin-28; Somatostatin precursor [Contains: Somatostatin Pecora; SGM SGM SGM SGM MAMMAILE SET SEADER (SAM SGM SGM SGM SGM SGM SGM SEQUENCE FROM N.A.

NCBI_TAXID=9940;

RN DELINB-99094691; PubMed-9880082;

RN DELINB-99094691; PubMed-9880082;

RM TLOCALIZATION of the preprosomatostatin-mRNA by in situ PRT SEQUELULAR LOCATION: Socreted.

TLOCALIZATION of the preprosomatostatin-mRNA by in situ PRT STRAIN-THE WEDLINGS TO THE SOMATOSTATIN FAMILY.

TO THE SWISS-PROT entry is copyright It is produced through the substantian sear on creating sear on the between the Swiss Institute of Bioinformatics and the Copyright of the Drochen Bioinformatics in Stitute. There are no creating sequires a license agreement (See http://www.ist. use by non-profit institutions as long as its content modified and this statement is not removed. Usage by modified and this statement is not removed.

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Best Local Similarity 48.8%; Pred. No. 0.1;
Matches 20; Conservative 4; Mismatches 14; Ind
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RESULT 10 SMS2_LOPAM

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                                                                                               uteleostei; Neoteleostei;
formes; Lophildae; Lophius
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Signal; Hydroxylation;
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                                                 [y10]somatostatin-14]
                                                                                                                                                                                                                                                                                                                              instational processing of preprosomatostatin-II examined to atom bombardment mass spectrometry."; Chem. 262:12692-12699(1987).
                                                                            Euteleostomi
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of cDNAs encoding two dis
In the endocrine pancreas
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5E14605D7B9A46FE CRC64;
                                                                                  Vertebrata;
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pair of basic residues; Hormone;
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                                                                                                                                                                                                                                                                                                                    J.E.;
                                                                                                                                                                                                                                                                                    PARTIAL SEQUENCE, AND HYDROXYLATION. MEDLINE-87308304; PubMed-2887572; Andrews P.C., Nichols R., Dixon J.E. "Post-translational processing of pr
                 Last sequence
                                                 [Contains
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                                                                                               Neopterygii; Tele
Paracanthopterygii
                                                                                                                                                                          PubMed-6107860;
                                                                                                                                                                                     Hobart P.M., Crawford R., Shen L. "Cloning and sequence analysis of somatostatin precursors found in
                                                                                                                                                                                                                                                                                                                                                                                                              SOMATOSTATIN
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uropean Bioinformatics
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                                                                                                                                                                                                                                                                                                                                                                                               SUBCELLULAR LOCATION: MISCELLANEOUS: SOMATO
                                               precursor
                                                                                                                                                                                                                                   anglerfish.";
Nature 288:137-141(1980)
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                                                            americanus
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22; Conser
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01-OCT-1989 (Rel
16-OCT-2001 (Rel
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                                                                                            Actinopterygii;
Acanthomorpha;
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                                                 Somatostatin
                                                                 Lophius ame
Eukaryota;
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P26917;
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SMS_BOVIN
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PubMed-9163424
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Endothelial nitric oxide
haracterization of a dist
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                                Freiberg C.A., Fellay
Perret X.;
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NCBI_TaxID=9913;
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Nitric-oxide s
III) (NOSIII)
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutel
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora;
Bovidae; Bovinae; Bos.
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Bacteria; Proteobacteria; alpha subdivision; Rhizobiaceae
Rhizobiaceae; Rhizobium.
NCBI_TaxID=394;
01-AUG-1992 (Rel. 23, Created)
01-AUG-1992 (Rel. 23, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Somatostatin precursor (Contains: Somatostatin-28; Somates
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                                                                                                                                              37; PubMed=2899837;
J.W., Li W.H., Luo C.C., Frazier M.L.,
                                                                                                                                                                           "Structure and evolution of somatostatin genes."; Mol. Endocrinol. 2:209-216(1988).
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RHISN
Y4RB_RHISN
STANDARD;
P55635;
01-NOV-1997 (Rel. 35, Created)
01-NOV-1997 (Rel. 35, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Putative integrase/recombinase Y4RB.
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7; Mismatches
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Best Local Similarity 45.2%;
Matches 19; Conservative
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netical protein; DNA recombination; DNA integration;
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MEDLINE-93055452; Pubmec.
Nishida K., Harrison D.G., Navas c.
"Molecular cloning and characterization of the cc
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Pred. No. 2.3;
1; Mismatches
                                                                                                                                   ween the Swiss Institute of Bloinform
European Bloinformatics Institute. The by non-profit institutions as long
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                                                                                                                                                                                             modified and this statement is not remorentities requires a license agreement (or send an email to license@isb-sib.ch)
                       Perret X.; "Molecular basis of symbiosis between Nature 387:394-401(1997).
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C., Harrison J.K., Barber C
D.D., Lynch K.R., Peach M.
Ir cloning and expression c
                                                                                                                    entry is copyright iss Institute of Bio
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                                  J. Biol. Chem. 267:15274-15276(1992).
[4]
MYRISTOYLATION.
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Busconi L., Michel T.;
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[5]
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MEDLINE-21552923; Pubmed-11695891;

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Elochemistry 40:13448-13455(2001).

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X RAMDLINE-21336567; Pubmed-11331290;

M. Masters B.S.S., Poulos T.L.;

Raman C.S., Li H., Martasek P.; Babu B.R., Griffith O.W., S. Masters B.S.S., Poulos T.L.;

Implications for isoform-selective inhibitor design derive Dinding mode of bulky isothioureas to the heme domain of en intric-oxide synthase.";

J. Biol. Chem. 276:26486-26491(2001)

L. J. Biol. Chem. 276:26486-26491(2001)

T. SANGDUCTION: PRODUCES NITRIC OXIDE (NO) WHICH IS IMPLICATE FUNCTION: PRODUCES NITRIC OXIDE (NO) WHICH IS IMPLICATE FORCED ANGIOGENESIS IN CORONARY VESSELS FOUNCED NOTIVE THOUGH TER ACTIVATION OF PLATE FORCED SECONDERS BLOOD CLOTTING THROUGH THE ACTIVATION OF PLATE CATALYTIC ACTIVITY: L-arginine + N NADPH + M O(2) = cit nitric oxide + N NADPP + H O(2) = cit nitric oxide + N NADPP + H O(2) = cit nitric oxide + N NADPP + M O(2) = cit nitric oxide + N NADPP + M O(2) = cit nitric oxide + N NADPP + M O(2) = cit nitric oxide + N NADPP + M O(2) = cit nitric oxide + N NADPP + M O(2) = cit nitric oxide + N NADPP + M O(2) = cit nitric oxide + N NADPP + M O(2) = cit nitric oxide + N NADPP + M O(2) = cit nitric oxide + N NADPP + M O(2) = cit nitric oxide + N NADPP + M O(2) = cit nitric oxide + N NADPP + N O(2) = cit nitric oxide + N NADPP + N O(2) = cit nitric oxide + N NADPP + N O(2) = cit nitric oxide + N NADPP + N O(2) = cit nitric oxide + N NADPP + N O(2) = cit nitric oxide + N NADPP + N O(2) = cit nitric oxide + N NADPP + N O(2) = cit nitric oxide + N NADPP + N O(2) = cit nitric oxide + N NADPP + N O(2) = cit nitric oxide + N NADPP + N O(2) = cit nitric oxide + N NADPP + N O(2) = cit nitric oxide + N NADPP + N O(2) = cit nitric oxide + N NADPP + N O(2) = cit nitric oxide + N NADP + N O(2) = cit nitric 
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MEDLINE-96102197; PubMed-8524847;
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Rageting.";
Proc. Natl. Acad. Sci. U.S.A. 92:11776-11780(1995).
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Raman C.S., Li H., Martasek P., Kral V., Masters B.S.S., Raman C.S., Li H., Martasek P., Kral V., Masters B.S.S., Raman C.S., Martasek P., Kral V., Masters B.S.S., Li H., Raman C.S., Martasek P., Kral V., Masters B.S.S., Li H., Raman C.S., Martasek P., Kral V., Masters B.S.S., Li H., Raman C.S., Martasek P., Kral V., Masters B.S.S., Li H., Raman C.S., Martasek P., Kral V., Masters B.S.S., Li H., Raman C.S., Martasek P., Kral V., Masters B.S.S., Li H., Raman C.S., Martasek P., Kral V., Masters B.S.S., Li H., Raman C.S., Martasek P., Kral V., Masters B.S.S., Li H., Raman C.S., Martasek P., Kral V., Masters B.S.S., Li H., Raman C.S., Martasek P., Kral V., Masters B.S.S., Li H., Raman C.S., Martasek P., Kral V., Masters B.S.S., Li H., Raman C.S., Martasek P., Kral V., Masters B.S.S., Li H., Raman C.S., Martasek P., Kral V., Masters B.S.S., Li H., Raman C.S., Martasek P., Kral V., Masters B.S.S., Li H., Raman C.S., Martasek P., Kral V., Masters B.S.S., Li H., Raman C.S., Martasek P., Kral V., Masters B.S.S., Li H., Raman C.S., Martasek P., Kral V., Masters B.S.S., Li H., Raman C.S., Martasek P., Kral V., Masters B.S.S., Li H., Raman C.S., Martasek P., Kral V., Masters B.S.S., Li H., Raman C.S., Martasek P., Raman C.S., Martasek P.,
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[8]

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oxide synthase.";
.. Chem. 267:15274-15276(1992)
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369; FLAVODOXIN.
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Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Saccharomyces.
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SEQUENCE FROM N.A.
STRAIN-NRRL'Y-12624;
COSTANZO M.C., FOX T.D.;
COSTANZO M.C., FOX T.D.;
Submitted (SEP-1997) to the EMBL/GenBank/DDBJ databases.
-!- FUNCTION: REQUIRED FOR EXPRESSION OF THE MITOCHONDRIAL GENCYTOCHROME C OXIDASE SUBUNIT III (COX3). PET122 SEEMS TO IDIRECTLY INTERACTING WITH THE SMALL RIBOSOMAL SUBUNIT TO ITRANSLATION INITIATION ON THE COXIII MRNA (BY SIMILARITY)
-!- SUBCELLULAR LOCATION: MITOCHONDRIAL; INNER MEMBRANE ASSOCTING WILLARITY).
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80AFEC2A7238262F CRC64;
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Pred. No. 2:8;
10; Mismatches
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15-JUL-1998 (Rel. 36, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
PET122 protein, mitochondrial precursor.
PET122.
Score 67; DB Pred. No. 10; 4; Mismatches
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Similarity 27.0%;
17; Conservative 1
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P01167;
21-JUL-1986
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Lo-UCT-2001 (Rel. 40, Last annotation update)
Somatostatin precursor [Contains: Antrin; Somatostatin-28;
Somatostatin-14].
SST OR SMST.
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Montminy M.R., Goodman R.H., Horovitch S.J., Habener J.F.
"Primary structure of the gene encoding rat preprosomatos
Proc. Natl. Acad. Sci. U.S.A. 81:3337-3340(1984).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE-85006903; PubMed-6148343;
Tavianini M.A., Hayes T.E., Magazin M.D., Minth C.D., Dixelation, characterization, and DNA sequence of the rat
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Goodman R.H., Jacobs J.W., Dee P.C., Habener J.F.;
"Somatostatin-28 encoded in a cloned cDNA obtained from
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Sciurognathi; Muridae;
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dontminy M.R., Low M.J., H
of rat preprosomatostatin.
Biol. 188:31-47(1985).
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1 D.C., Roos B.A.;
Statin. Structure
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Taylor W.L., Minth C.D., I
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J. Biol. Chem. 258:8788-8793(1983)
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Fuhrmann G., Heilig R., Kempf J.,
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EMBL: NO1218, AAA42167.1;

EMBL: NO1787, AAA42167.1;

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EMBL: S08416; CAA35831.1;

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EMBL: S08416; CAA5581.1;
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Search completed: March 21, 2003, 11:38:40 Job time: 13.3348 secs

	GenCore version 5.1.4_p5_4578 Copyright (c) 1993 - 2003 Compugen Ltd.
OM protein - pro	protein search, using sw model
Run on:	March 21, 2003, 11:38:56; Search time 18.4549 Seconds (without alignments) 447.986 Million cell updates/sec
Title: Perfect score: Sequence:	US-09-727-739B-17 450 1 MRVSQIHCALALLGLALAICRWRPRKVKGPQLKAKEDLER 86
Scoring table:	BLOSUM62 Gapop 10.0 , Gapext 0.5.
Searched:	283224 seqs, 96134422 residues
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C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999.#text_change 10-Sep-1999 C;Date: 220630 C;Accession: S20630 B;Nata, K.; Kobayashi, T.; Karahashi, K.; Kato, S.; Yamamoto, H.; Yonekura, H.; Okasubmitted to the EMBL Data Library, June 1991 A;Description: Nucleotide sequence determination of chicken somatostatin precursor A; Reference number: S20630 A;Reference number: S20630 A;Status: preliminary A;Molecule type: mRNA A;Residues: 1-116 <NAT> A;Residues: 1-116 <NAT> A;Residues: 1-116 <NAT> A;Residues: 1-116 <NAT> A;Residues: C;Superfamily: somatostatin

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bolleau, G.;
11-2916, 1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Johnsen, A.H
303, 263-268
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
29; Conser
    Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               -108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    P.; F.
                     21;
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Best Local
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                     pancreatic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PID:964018
                                                                                                                                                                                                       RIDES
Somatostatin-14 precursor - channel catfish
Nalternate names: somatostatin I
N'Alternate names: somatostatin I
C'Species: Ictalurus punctatus (channel catfish)
C'Date: 30-Jun-1900 # Seaguence_revision 31-Dec-1993 #text_change 18-Jun
C'Accession: 500292; A93877; A92347; A94435
N'AITLE: The structure of Cloned DNA complementary to catfish pancreat.
A'Reference number: 500292; MUID: 82265698; PMID: 6179939
A'Accession: 500292; MUID: 82265698; PMID: 6179939
A'Accession: 500292; MUID: 8265698; PMID: 6179939
A'Accession: A. Colliter, K.J.; Deschenes, R.J.; Wetth, H.L.; Dixon, J.J.
Cross-references: EMBL: V00607; NID: 964017; PIDN: CAA23877.1; PID: 9640
A'TATJOY, W.L.; Colliter, K.J.; Deschenes, R.J.; Wetth, H.L.; Dixon, J.J.
A'TATJOY, W.L.; Colliter, R.J.; Deschenes, R.J.; Wetth, H.L.; Dixon, J.J.
A'TILE: Sequence analysis of a cDNA coding for a pancreatic precursor A; Reference number: A3897; MUID: 82082515; PMID: 617821
A'Accession: 260: 88170994
A'Accession: 260: 88170994
A'Accession: 260: 88170994
A'Accession: 260: 88170944
A'
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C; Species: Ictalurus punctatus (channel catfish)
C; Date: 13-Sep-1996 #sequence_revision 13-Sep-1996 #text_change
C; Accession: I50798
R; Dixon, J.E.; Andrews, P.C.
Adv. Exp. Med. Biol. 188, 19-29, 1985
A; Title: Somatostatins of the channel catfish.
A; Reference number: I50798; MUID: 85303576; PMID: 2863931
A; Accession: I50798
A; Status: preliminary; translated from GB/EMBL/DDBJ
A; Molecule type: mRNA
A; Residues: 1-114 <DIX>
A; Residues: 1-114 <DIX>
C; Superfemily: somatostatin
C; Superfamily: somatostatin
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S REQCALALESIALAVGTVSAAPSDPRE
                                                             64 PCLRWRPRKVKGPQLKAKEDLER 86
| | : | | | | 5
56 --LLSEPSQTENEALES-EDLSR 75
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1. Acad. Sci. U.S.A. 82, 277-281, 1985
Processing of an anglerfish somatostatin precursor
Ice number: A94038; MUID:85113184; PMID:2857489
                                                                                                                                                                                                                                                                                                       references: GB:V00641; GB:J00947; GB:M23199; NID imental source: islet tissue (endocrine pancreas)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   J.; White, J.W.; Li, W.H.; Luo, C.C.; Frazier, Scrinol. 2, 209-216, 1988
Structure and evolution of somatostatin genes.
Structure A40929; MUID:88288237; PMID:2899837
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     somatostatin
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Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            inhibits the release
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          pyrrolidone carboxylic a bonds: #status experimen
                                                                                                                                                          t, P.; Crawford, R.; Shen, L.; Picte 288, 137-141, 1980
E. Cloning and sequence analysis of cence number: A93236; MUID:81052423;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chem. 262, 12692-12699, 1987
Post-translational processing of pree number: A27376; MUID:87308304;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (SS-14); somate taurus (cattle)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    neuropeptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score
Pred.
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Species: Bos primigenius taurus (catt
Date: 06-Mar-1992 #sequence_revision
                                                                                                              #sequence_revision
A94038; A27376; A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         -89, 'G', 91-125
                                                                                                                                                        Shen,
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36.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    hydroxylysine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  type: protein: 98-125 <SPI>
                                                                                                                                                                                                                                                                                                                                                         Noe, B.D.
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22; Conser
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NiAlternate names: PSS1 protein
C; Species: Rana ridibunda (laughing frog)
C; Date: 11-Apr-1997 #sequence_revision 09-May-1997 #text_change 16-Jul-1999
C; Accession: JC6166
R; Tostivint, H.; Lihrmann, I.; Bucharles, C.; Vieau, D.; Coulouarn, Y.; Fournie: Proc. Natl. Acad. Sci. U.S.A. 93, 12605-12610, 1996
A; Title: Occurrence of two somatostatin variants in the frog brain: Characteriz: A; Reference number: JC6166; MUID: 97057290; PMID: 8901629
A; Contents: brain
A; Accession: JC6166
A; Molecule type: mRNA
A; Residues: 1-115 < TOS>
A; Consorted: This protein acts both as a neurotransmitter/neuromodulator and a ho; C; Genetics:
A; Genetics:
A; Genetics:
C; Superfamily: somatostatin
C; Superfamily: brain; hormone
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                                                                                                  RESULT 6
A28968
Somatostatin I precursor - crab-eating macaque
N;Alternate names: preprosomatostatin
N;Contains: somatostatin 14 (SS-14); somatostatin 28 (SS-28)
C;Species: Macaca fascicularis (crab-eating macaque)
C;Date: 30-Jun-1989 #sequence_revision 31-Jan-1997 #text_chan
C;Accession: A28968
R;Travis, G.H.; Sutcliffe, J.G.
Proc. Natl. Acad. Sci. U.S.A. 85, 1696-1700, 1988
A;Title: Phenol emulsion-enhanced DNA-driven subtractive cDNA
A;Reference number: A28968; MUID:88144503; PMID:2894033
A;Accession: A28968
A;Molecule type: mRNA
A;Residues: 1-116 <TRA>
A;Residues: 1-116 <TRA>
A;Coss-references: GB:M19318; NID:g342298; PIDN:AAA36908.1;
C;Comment: Somatostatin
C;Keywords: hormone; neuropeptide
F;1-24/Domain: signal sequence #status predicted <SIG>F;25-88/Domain: signal sequence #status predicted <M18>
F;25-88/Domain: propeptide #status predicted <M18>
F;103-116/Product: somatostatin-18 #status predicted <M14>
F;105-116/Disulfide bonds: #status predicted
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   S RLQCALAALSIVLALGCVTGAPSDPRL----RQFLQKSLAAA----AG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28;
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Pred. No. 0.22;
; Mismatches 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 79.5; DB; Pred. No. 0.1; 13; Mismatches
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                           --LLSEPNOTENDALE-PEDLSO
PCLRWRPRKVKGPQLKAKEDLER
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Local Similarity 47.8%;
les 22; Conservative 6
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| Similarity 34.9%;
| 29; Conservative
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fast

preprosomatostatin-II PMID:2887572

Gaps

15;

17;

experimental

#status

<MAT>

<SIG>

<PRO>

pyroglutamic.

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Length

--PDLDLASRR--LLQRALAAALPHRSGVSER

Chan,

31-Jan-1997 #text_change 18-Jun-1999

(88-58)

M. L.;

of

hydroxylysine-cont

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PIDN: CAA23987

NID:964030;

change

1 #text_cha 81; A93236 Rutter, W.

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(strain PAO1)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           anome sequence of Pseudomonas aeruginosa PA01, A82950; MUID:20437337; PMID:10984043
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     aeruginosa
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                                                                                                                                                                                                                                                                                                                                                  K.R.;
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                                                                                                                                                     subtilisin
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Pred. No. 2.7;
8; Mismatches
                                                                                                                                                ag12;
                                                                                                                                                                                                     Score 68; DB Pred. No. 12:
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Mismatches
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                                                                                                                                                     proteinase
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Similarity 33.9%;
1; Conservative
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                                                                                                                                                                                                                                    Similarity 28.3
18; Conservative
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Y.; Brody,
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erence number: A8299
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d to GenBank,
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amas, S.; Marsden
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Conserved hypothetical protein XF2472.[imported] - Xylella fastidiose Cispecies: Xylella fastidiosa
C;Species: Xylella fastidiosa
C;Species: Xylella fastidiosa
C;Species: Yylella fastidiosa
C;Accession: A82554
R;Anonymous, The Xylella fastidiosa Consortium of the Organization fature 406, 151-157, 2000
Title: The genome, sequence of the plant pathogen Xylella fastidios Reference number: A82515; MuID:20365717; PMID:10910347
A;Note: for a complete list of authors see reference number A59328 by Accession: A82554
A;Note: for a complete list of authors see reference number A59328 by Accession: A82554
A;Status: preliminary
A;Note: for a complete list of authors see reference number A59328 by Accession: A82554
A;Status: preliminary
A;Residues: 1-435 <SIM
A;Residues: 1-435 <SIM
A;Residues: 1-435 <SIM
A;Residues: 1-435 <SIM
A;Residues: C; Bl-Dorry, H; Facincani, F.A.; Ferreira, A;Brinosa, M. R.S.; Bueno, M. R.P.; Buenco, M. R.P.; Buenco, M. R.P.; Ferreira, A. Brinosa, M. R.S.; Bueno, M. R.P.; Buenco, M. A.; Kemper, E.L.; Kitajima, J.P.; Krieger, J. E., Chado, M. A.; Madelra, A. M. B. N.; Madelra, H. M.F.; Mathors: Martins, B. M.F.; Matsukuma, A. Y.; Menck, C.F. M.; Miracca, F. G.; Nunes, L. R.; Oliveira, M. A.; Ge Oliveira, M. C.; de Oliveira, R. R. Silva, A. Silva, A.
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T51335
subtilisin-like proteinase AIR3, auxin-induced, [imported] - Arabidog subtilisin-like proteinase AIR3, auxin-induced, [imported] - Arabidog subtilisin-like proteinase AIR3, auxin-induced, [imported] - Arabidog C; Species: Arabidopsis thaliana (mouse-ear cress)
C; Date: 18-Aug-2000 #sequence_revision 18-Aug-2000 #text_change 03-PC; Accession: T51335
R; Neuteboom, L.W.; Ng, J.M.Y.; Kuyper, M.; Clijdesdale, O.R.; Hooyka Plant Mol. Biol. 39, 273-287, 1999
A; Title: Isolation and characterization of cDNA clones corresponding A; Reference number: 225377; MUID:99178779; PMID:10080694
A; Accession: T51335
A; Accession: T51335
A; Accession: T51335
A; Molecule type: mRNA
A; Molecule type: mRNA
A; Residues: 1-758 < NEU>
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C;Superfamily: Aquifex aeolicus N-ethylammeline chlorohydrolase
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F;1-24/Domain: signal sequence #status predicted <SIG
F;25-88/Domain: propeptide #status predicted <PRO>
F;89-116/Product: somatostatin-28 #status predicted <
F;103-116/Product: somatostatin-14 #status predicted
F;105-116/Disulfide bonds: #status predicted
                                                                                                                                                                       DB 1;
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Best Local Similarity 45.2%; Pred. No. 1.8;
Matches 19; Conservative 7; Mismatches
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of

PID:97

NID:97227332; PIDN:AAF42400.1;

Neisseria menigitidis

Length 259;

Indels

Mismatches

Score

Pred

Klee, M.A.;

er, C.; Quail,

Churcher,

S.D.; Chu Mungall,

Bentley,

S.; Moule,

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GB:AL157959; NID:g7378778; PIDN:CAB83652.1;
A, Strain Z2491
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bmitted to the Protein Sequence Database,
Reference number: 222516
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                exodeoxyribonuclease III
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larity 29.0%;
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PID:9163427
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, V.; Pizza, M.
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AAccession: A46033
AAccession: A46034
AAccession: A46034
AAccession: A6031
AAccession: A6045 (17,862-1205 c.LAM2)
AACCESS-TefeTences: G81809525; NID:165976
AACCESS-TefeTences: G81809525; NID:165976
AAACCESSION: ABATISON, D.G.; NAVAS, J.P.; Fisher, A.A.; Dockery, S.P.; Ne RAISHIDIAGA K.Y.; Harrison, D.G.; NAVAS, J.P.; Fisher, A.A.; Dockery, S.P.; Ne AAACCESSION: 145945; MUID:9305452; PRID:1385480
AAACCESSION: 145945; MUID:93055452; PRID:1385480
AAACCESSION: 145945; MUID:93055452; PRID:1385480
AAACCESSION: A15945; MUID:93055452; PRID:1385480
AAACCESSION: A15945; MUID:93055452; PRID:1385480
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R; Tettelin, H.; Saunders, N.J.; Heidelberg, J.; Jeffries, A.C.; Nels Hickey, E.K.; Haft, D.H.; Salzberg, S.L.; White, O.; Fleischmann, R. ri, H.; Oin, H.; Vamathevan, J.; Gill, J.; Scarlato, V.; Masignani, Science 287, 1809-1815, 2000
A; Authors: Grandi, G.; Sun, L.; Smith, H.O.; Fraser, C.M.; Moxon, E. A; Authors: Grandi, G.; Sun, L.; Smith, H.O.; Fraser, C.M.; Moxon, E. A; Title: Complete genome sequence of Neisseria meningitidis serogrou A; Reference number: A81000; MUID:20175755; PMID:10710307
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14.9%; Score 67; DB 1; Length
Best Local Similarity 35.5%; Pred. No. 25;
Matches 27; Conservative 4; Mismatches 25; Inde
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score:

Title: Perfect sc Seguence:

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Run on:

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Total number

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1: /cgn2_6/ptodata/1/pubpaa/US08_NEW_PUB.pep:
2: /cgn2_6/ptodata/1/pubpaa/PCT_NEW_PUB.pep:
3: /cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep.
4: /cgn2_6/ptodata/1/pubpaa/US07_NEW_PUB.pep.
5: /cgn2_6/ptodata/1/pubpaa/US07_NEW_PUB.pep.
6: /cgn2_6/ptodata/1/pubpaa/US07_NEW_PUB.pep.
7: /cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep.
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13: /cgn2_6/ptodata/1/pubpaa/US60_NEW_PUB.pep.
14: /cgn2_6/ptodata/1/pubpaa/US60_NEW_PUB.pep.
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5.1.4_p5_457
Compugen Ltd
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10 US-09-833-745-63

9 US-09-984-271-131

10 US-09-766-396-3

12 US-10-062-375-3

9 US-10-062-375-3

9 US-10-001-054-42

9 US-10-028-072-132

9 US-10-121-049-132

9 US-10-123-904-132

9 US-10-123-904-132

9 US-10-176-918-132

9 US-10-176-918-132

9 US-10-176-918-132

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9 US-10-142-431-132
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Maximum Match 100%
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Query Match

Score

Pred

657.5 662.5 661.5

122 100 110 112 113 114 116

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US-09-833-745-63
Sequence 63, Application US/09833745
Sequence 63, Application US/09833745
Sequence 63, Application US/09833745
Sequence 63, Application US/09833745
Sequence 63, Application:
GENERAL INFORMATION:
APPLICANT: ROBERTS, JOSEPH
APPLICANT: SETHURAMAN, NATARAJAN
APPLICANT: MACALLISTER, THOMAS
TITLE OF INVENTION: BIOACTIVE HISTIDINE AMMONIA LYASE
FILE REFERENCE: 078728/0106
CURRENT APPLICATION NUMBER: 60/197,770
PRIOR FILING DATE: 2000-04-14
NUMBER OF SEQ ID NOS: 66
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 63
LENGTH: 513
MANDE: DATE
MANDE
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N: EXPRESSED IN FETAL LIVER, SIGNAL = 2.6
N: EXPRESSED IN PLACENTA, SIGNAL = 2.6
N: EXPRESSED IN ADULT LIVER, SIGNAL = 2.7
N: EXPRESSED IN BRAIN, SIGNAL = 2.7
N: EXPRESSED IN HEART, SIGNAL = 2.5
N: EXPRESSED IN HEART, SIGNAL = 2.5
N: EXPRESSED IN HEART, SIGNAL = 2.5
N: EXPRESSED IN HELA, SIGNAL = 2.5
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Pred. No. 0.81;
11; Mismatches 20;
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Best Local Similarity 38.2%; Pred. No. 9.5;
Matches 21; Conservative 5; Mismatches
PRIOR APPLICATION NUMBER: PCT/US01/00670
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 09/608,408
PRIOR FILING DATE: 2000-06-30
PRIOR FILING DATE: 2001-01-29
NUMBER OF SEQ ID NOS: 49117
SOFTWARE: Annomax Sequence Listing Engine ve; SeQ ID NO 41361
LENGTH: 95
TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
OTHER INFORMATION: EXPRESSED IN PLACENTA, SOTHER INFORMATION: EXPRESSED IN BRAIN, SIGN, OTHER INFORMATION: EXPRESSED IN BONE MARROW OTHER INFORMATION: EXPRESSED IN HEART, SIGN, OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL OTHER INFORMATION: EXPRESSED IN HELA, SI
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Similarity 27.7%;
23; Conservative
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US-09-984-271-131
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Human Secreted Proteins
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Pred. No. 9.9;
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                          US/09/984,271
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Patent No. US20020013456A1
GENERAL INFORMATION:
APPLICANT: Sutcliffe, Gregor
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19; Conservative
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                                                                                      CATION NUMBER:
                                                                                                                                                                                                                                                                                    M: Homo sapiens
                                                                                                                                                                                                         F. SEQ ID NOS: Patentin Va
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9-984-271-131
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US-09-766-396-3
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                                 CALALLGLALAICS-QGAASQPDLDLASRRLLQRALAAA
   Mismatches
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: 2001-11-30
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KOLODNER, Richard
                                                                                                                                         Application US/09470276 to. US20020187469A1
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o. US20020192209A1
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1998-07-02
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0; Conservative
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                                                                                                                                                                       FORMATION
                                                                                                                                                                                                                                      INVENTION
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US-09-470-276-54
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19;
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Best Local S
Matches 30
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/062,375
FILING DATE: 30-Jan-2002
CLASSIFICATION: <Unknown>
PRIOR APPLICATION NUMBER: US/08/857,389
FILING DATE: <Unknown>
APPLICATION NUMBER: US/08/857,389
FILING DATE: <Unknown>
REGISTRATION NUMBER: 31,796
REGISTRATION NUMBER: 31,796
REFERENCE/DOCKET NUMBER: 22908-0002
TELECOMMUNICATION:
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Siggins, George R.
Henriksen, Steven J.
Henriksen, CORTISTATIN: NEUROPEPTIDES,
COMPOSITIONS AND METHODS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 26

CORRESPONDENCE ADDRESS:
ADDRESSEE: THE SCRIPPS RESEARCH INSTITUTE
STREET: 10666 No. US20020133000A1th Torrey
CITY: La Jolla
STATE: California
                                                                                                                                                                                                                                                   10;
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                                                                                                                                                                                                                                                                                                 2 CALAALCIVLALGGVTGAPSDPRL----RQFLQKSLAAA
                                                                                                                                                                                                                                                                                8 CALALLGLALAICS-QGAASQPDLDLASRRLLQRALAAA
                                                                                                                                                                                                                th 13.9%; Score 62.5; Distinction 48.7%; Pred. No. 3.7; 19; Conservative 5; Mismatches
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        TELECOMMUNICATION INFORMATION:

TELEFAX: (415) 324-7041

TELEFAX: (415) 324-7041

TELEFAX: (415) 324-0638

SEQUENCE CHARACTERISTICS:

LENGTH: 110 amino acids

TYPE: amino acid

TOPOLOGY: linear

MOLECULE TYPE: protein

FRAGMENT TYPE: C-terminal

SEQUENCE DESCRIPTION: SEQ ID NO: 3:

US-09-766-396-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear

MOLECULE TYPE: protein

FRAGMENT TYPE: C-terminal

SEQUENCE DESCRIPTION: SEQ ID NO: 3:
US-10-062-375-3
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                   RESULT 5
US-10-062-375-3
; Sequence 3, Application US/10062375
; Patent No. US20020133000A1
; GENERAL INFORMATION:
    APPLICANT: Sutcliffe; Gregor J
de Lecea, Luis
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de Lecea, Luis
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(415) 324-0638
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCE CHARACTERISTICS:
LENGTH: 110 amino acids
TYPE: amino acid
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SEQUENCE CHARAC
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PRIOR APPLICATION NUMBER: 60/08345
PRIOR PLING OARE: 1998 0-62
PRIOR PELING OARE: 1998 0-62
PRIOR PELING OARE: 1998 0-62
PRIOR PAPLICATION NUMBER: 60/0869
PRIOR PELING DATE: 1998 0-62
PRIOR PELING DATE: 1999 0-63
PRIOR

UMBER: PCT/US99/20594 1999-09-08 UMBER: PCT/US99/28313 1999-11-30 UMBER: PCT/US00/14941 2000-05-30 UMBER: PCT/US00/15264 2000-06-02 NUMBER: PCT/US99/00106 1999-01-05 NUMBER: PCT/US99/05028 1999-03-08 R: PCT/US99/20111 -09-01 PCT/US99/28634 ER: PCT/US00/00376 0-01-06 MBER: PCT/US99/08615 999-04-20 3600E/66SD/ PCT/US00/03565 IBER: PCT/US00/13705 PCT/US99/12252 PCT/US99/30999 MBER: PCT/US00/04341 000-02-18 MBER: PCT/US00/14042 000-05-22 ER: PCT/US00/05841 0-03-02 /US98/18824 PCT/US99/28551 PCT/US00/06884 /US00/08439 омыЕR: PCT/US00/3087: 2000-11-10 946374 872035 882636 918585 924419 929404 SR: PCT/)-03-30 001-08-06 JEER: 0970 JMBER: PCT, 1999-06-02. JMBER: PCT, 9-12-20 98-09-10 -12-16 -02 - 18-03-15 -07-30 2001-09-04 NUMBER: PCT -12 - 02-08-06 -08-28 -02-11 ER: 09/ -05-MBER: IMBER: MBER: MBER: MBER: NUMBER: NUMBER: NUMBER: CATION PRIOR PRIOR RIOR RIOR RIOR IOR IOR FOR OR RIOR RIOR RIOR RIOR IOR IOR COR RIOR RIOR RIOR RIOR RIOR RIOR RIOR RIOR RIOR RIOR

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Pred. No. 17;
6; Mismatches
PRIOR APPLICATION NUMBER: PCT/US00/32678

PRIOR FILING DATE: 2000-12-01

PRIOR APPLICATION NUMBER: PCT/US01/06520

PRIOR FILING DATE: 2001-02-28

PRIOR APPLICATION NUMBER: PCT/US01/17092

PRIOR FILING DATE: 2001-03-01

PRIOR FILING DATE: 2001-05-25

PRIOR FILING DATE: 2001-06-01

PRIOR FILING DATE: 2001-06-01

PRIOR FILING DATE: 2001-06-20

PRIOR FILING DATE: 2001-06-29

PRIOR FILING DATE: 2001-08-29

NUMBER OF SEQ ID NOS: 91

SEQ ID NO 42
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US-10-028-072-132
; Sequence 132, Application US/10028072
; Publication No. US20030004311A1
; GENERAL INFORMATION:
; APPLICANT: Baker, Kevin P.
; APPLICANT: Beresini, Maureen
; APPLICANT: DeForge, Laura
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   R APPLICATION NUMBER: 60/056974
R FILING DATE: 1997-08-26
R APPLICATION NUMBER: 60/059113
R FILING DATE: 1997-09-17
R APPLICATION NUMBER: 60/059115
R FILING DATE: 1997-09-17
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7-09-17
ER: 60/059122
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60/059184
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Filvaroff, Ellen
Gao, Wel-Qiang
Gerritsen, Mary E.
Goddard, Audrey
Godowski, Paul J.
Gurney, Austin L.
Sherwood, Steven
Smith, Victoria
Stewart, Timothy A.
Tumas, Daniel
                                                                                                                                                                                                                                                                                          ch 13.7%;
l Similarity 33.3%;
19; Conservative
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Wood, William
Zhang
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APPLICATION NUMBER:
FILING DATE: 1997-09
APPLICATION NUMBER:
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APPLICANT: Woo
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TITLE OF INVENT
FILE REFERENCE:
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R PELLING DATE: 1998-02-09
R FILING DATE: 1998-02-09
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R FILING DATE: 1998-03-20
R PELLING DATE: 1998-03-20
R FILING DATE: 1998-03-25
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R FILING DATE: 1998-03-27
R FILING DATE: 1998-04-09
R FILING DATE: 1998-04-09
R FILING DATE: 1998-04-09
R FILING DATE: 1998-04-15
R APPLICATION NUMBER: 60/081817
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R APPLICATION NUMBER: 60/081818
R FILING DATE: 1998-04-15
R APPLICATION NUMBER: 60/081818
R FILING DATE: 1998-04-24
R APPLICATION NUMBER: 60/081818
R FILING DATE: 1998-04-24
R APPLICATION NUMBER: 60/081818
R FILING DATE: 1998-05-07
R APPLICATION NUMBER: 60/081818
R FILING DATE: 1998-05-07
R APPLICATION NUMBER: 60/081338
R FILING DATE: 1998-05-07
R APPLICATION NUMBER: 60/085318
R FILING DATE: 1998-05-07
R APPLICATION NUMBER: 60/085318
R FILING DATE: 1998-05-13
R APPLICATION NUMBER: 60/085318
R FILING DATE: 1998-05-13
R APPLICATION NUMBER: 60/085579
R FILING DATE: 1998-05-15
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R APPLICATION NUMBER: 60/089532
R FILING DATE: 1998-06-17
R APPLICATION NUMBER: 60/089599
R FILING DATE: 1998-06-17
R APPLICATION NUMBER: 60/089907
R FILING DATE: 1998-06-18
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3R: 60/0
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2R: 60/C
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TION NUMBER: 60/0
DATE: 19/98-06-1
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18 AICSQ-GAASQP-DLDLASRRLLQRALAAALPHRSGVSERWRTFY

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Length 333;

DB 9;

Score 61.5; Pred. No. 1

9

similarity 33.3%; 19; Conservative

Query Match Best Local S Matches 19

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7091982

NUMBER: 60/0 : 1998-07-02 | NUMBER: 60/0 :: 1998-07-07

APPLICATION | FILING DATE:

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NUMBER: 60/0 1998-06-24 NUMBER: 60/0 1998-06-24 NUMBER: 60/0 NUMBER: 60/0 NUMBER: 60/0 NUMBER: 60/0 NUMBER: 60/0

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TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC NG THE SAME
        --VSDRYRFLYCYVPKVACSNWK 120
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Pred. No. 17;
6; Mismatches
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Application US/10121049
. US20030022239A1
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Similarity 33.3%;
19; Conservative
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Wood, William
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CURRENT APPLICATION NUMBER:
CURRENT FILING DATE: 2002-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
CRGANISM: Homo Sapien
US-10-121-049-132
                                                                         Sequence 132, Applipublication No. US2 GENERAL INFORMATION
                                                 RESULT 9
US-10-121-049-132
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LENGTH: 333
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Matches 19
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RESULT 10 US-10-123-904-132

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SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC ACIDS ENCODING THE SAME
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. US20030027275A1
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Similarity 33.3%;
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DeForge, Laura
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                                                                                                            ; TYPE: PRT
; ORGANISM: Homo Sapien
US-10-140-470-132
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US-10-140-470-132

Sequence 132, Application US/10140470

Publication No. US20030022331A1

GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
APPLICANT: Baker, Kevin P.
APPLICANT: Beresini, Maureen
APPLICANT: Beresini, Maureen
APPLICANT: Geovers. Luc
APPLICANT: Geovers. Luc
APPLICANT: Geovers. Luc
APPLICANT: Geovers, Lac
APPLICANT: Geovers, Lac
APPLICANT: Geovers, Lac
APPLICANT: Geovers, Lau J.
APPLICANT: Geovers, Lau J.
APPLICANT: Goodard, Audrey
APPLICANT: Goodard, Audrey
APPLICANT: Sherwood, Steven
APPLICANT: Sherwood, Steven
APPLICANT: Sherwood, Steven
APPLICANT: Sherwood, Steven
APPLICANT: Wood, William
APPLICANT: Wood, William
APPLICANT: Leven Through A.
APPLICANT: Connow, Daniel
APPLICANT: Connow, Conn
Sequence 132, Application US/10123904
Publication No. US20030022328A1
GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
APPLICANT: Beresini, Maureen
APPLICANT: Beresini, Maureen
APPLICANT: Desnoyers, Luc
APPLICANT: Geor, Wel-Qian
APPLICANT: Geor, Wel-Qian
APPLICANT: Georyers, Luc
APPLICANT: Georyers, Luc
APPLICANT: Georyers, Luc
APPLICANT: Georyers, Luc
APPLICANT: Godowski, Paul J.
APPLICANT: Godowski, Paul J.
APPLICANT: Gherey, Austin L.
APPLICANT: Sherwood, Steven
APPLICANT: Sherwood, Steven
APPLICANT: Tummas, Daniel
APPLICANT: Watanabe, Colin K
APPLICANT: Wata
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13.7%; Score 61.5; DB 9; Leng
Best Local Similarity 33.3%; Pred. No. 17;
Matches 19; Conservative 6; Mismatches 19; Ind
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; ORGANISM: Homo Sapien
US-10-123-904-132
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LicanT: Zhang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES A
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3330R1C288
CURRENT APPLICATION NUMBER: US/10/176,921
CURRENT FILING DATE: 2002-06-20
Prior Application removed - See File "
SEQ ID NO 132
LENGTH: 333
TYPE: PP"
TYPE: PP"
APPLICANT: Smith, Victoria
APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Watanabe, Colin K
APPLICANT: Watanabe, Colin K
APPLICANT: Wood, William
APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3330R1C382
CURRENT APPLICATION NUMBER: US/10/176,918
CURRENT FILING DATE: 2002-06-20
Prior Application removed - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 550
SEQ ID NO 132
LENGTH: 333
TYPE: PRT
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Pred. No. 17;
6; Mismatches
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5. US20030027276A1
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Beresini, Maureen
Beresini, Maureen
DeForge, Laura
Desnoyers, Luc
Filvaroff, Ellen
Gao, Wei-Qiang
Gerritsen, Mary E.
Goddard, Audrey
Goddard, Audrey
Godowski, Paul J.
Gurney, Austin L.
Sherwood, Steven
Smith, Victoria
Stewart, Timothy A.
Tumas, Daniel
Watanabe, Colin K
Wood, William
Zhang, Zamir
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Best Local Similarity 33.3%;
Matches 19; Conservative
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US-10-176-918-132
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US-10-176-921-132
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US-10-176-921-132

Sequence 132, Application No. US2003

GENERAL INFORMATION:
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Best Local S
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9-632-098-2

8-484-105-10

9-143-571-29

8-469-260A-344

8-176-427B-4

8-176-900C-9

8-460-900C-9

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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/
FILING DATE: 19920702
CLASSIFICATION: 435
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CHARACTERISTIC
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/note= "Potential proline directe
phosphorylation site"
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phosphorylation site"
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phosphorylation site"
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phosphorylation site"
                                                                                                                                                 /note= "FAD-Pyrophosphate binding domain"
                  "CA++/CAM binding domain"
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NAME/KEY: Binding-site
LOCATION: 937..947
OTHER INFORMATION: /note= "FAD-Isolalloxanthine OTHER INFORMATION: binding domain"
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NAME/KEY: Binding-site
LOCATION: 1012..1030
OTHER INFORMATION: /note- "NADPH-Ribose binding
FEATURE:
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LOCATION: 795..806
OTHER INFORMATION: /not
OTHER INFORMATION: dome
                                            NAME/KEY: Binding-site
LOCATION: 651..678
OTHER INFORMATION: /not
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LOCATION: 472..473
OTHER INFORMATION:
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OTHER INFORMATION:
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LOCATION: 33.34
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OTHER INFORMATION: 7
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LOCATION: 836..838
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 CALALLGLALAIC-SQGAAS---QPDLDLASRRLLQRALAAALPHRSGVSERWRTFYPNC
                                                                                                                                RESULT 2
US-08-319-866-10
; Sequence 10, Application US/08319866
; Patent No. 5929223
; GENERAL INFORMATION:
    APPLICANT: Tully, Timothy P.
    APPLICANT: Yin, Jerry C.
    APPLICANT: Regulski, Michael
    TITLE OF INVENTION: CLONING AND CHARACTERIZATION OF GENITILE OF INVENTION: ASSOCIATED WITH LONG-TERM MEMORY
    INVENESPONDENCE ADDRESS:
    ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C.
    STREET: Two Militia Drive
                                                                                                                                                                                                                                                                                                                                     ADDRESSEL.
STREET: Two Militia Dilv.
CITY: Lexington
STATE: Massachusetts
COUNTRY: USA
ZIP: 02173
COMPUTER READABLE FORM:
MEDIUM TYPE: FIDAPY disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/319,866
FILING DATE: 7-OCT-1994
CLASSIFICATION STATE: APPLICATION DATA:
APPLICATION NUMBER: 314
PRIOR APPLICATION DATA:
APPLICATION NUMBER: CSHL94-03
FILING DATE: ATTORNATION:
NAME: Granahan, Patricial
REGISTRATION NUMBER: GS14
FELEPHONE: (617) 861-9540
TELEPHONE: (617) 861-9540
TELEPHONE: (617) 861-9540
TELEPHONE: GS0 ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 1205 amino acids
TVPE: amino acid
                    --SRAPAPATPHAPI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length
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US-09-123-708-6
; Sequence 6, Application US/09123708
; Patent No. 6146887
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4;
                   Query Match
Best Local Similarity 35.5%;
Matches 27; Conservative
                                                                                 64 PCLRWRPRKVKGPQLK 79
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HYDROPHOBICALLY-MODIFIED PROTEIN COMPOSITIONS
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1999-06-03
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Local Similarity 26.2%;
es 17; Conservative
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larity 30.0%;
Conservative
                      NFORMATION:
                                                                        NUMBER:
                                                                                                                                                          V FOR SEQ ID NO:
CHARACTERISTICS
                                                                                                                      713-789-2679
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                                                     NUMBER
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US-09-325-256-24
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CVLWR 1162
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US-08-276-967-2
                                                                                                                                                         INFORMATION FOR
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TH: 396
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Best Local S
Matches 21
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Matches
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US-08-276-967-2
; Sequence 2, Application US/08276967
; Sequence 2, Application US/08276967
; Sequence 2, Application US/08276967
; Patent No. 5851817
GENERAL INFORMATION:
; APPLICANT: Garbers, David L.
; TITLE OF INVENTION: Specime TITLE OF INVENTION: Specime NUMBER OF SEQUENCES: 8
; CORRESPONDENCE ADDRESS:
; ADDRESSE: Arnold, White & Durkee STREET: P. O. Box 4433
                                                                                                                                                        Road, TPC-8
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/648,322
FILING DATE:
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Fitting, Thomas
REGISTRATION NUMBER: 34,163
REFERENCE/DOCKET NUMBER: 519.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Houston
STATE: Texas
COUNTRY: USA
ZIP: 77210-4433
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/276,967
FILING DATE: Submitted Herewith
Patent No. 6074872
GENERAL INFORMATION:
APPLICANT: Sutcliffe, Gregor J.
APPLICANT: de Lecea, Luis
TITLE OF INVENTION: CORTISTATIN: NEUROPEPTIDES,
TITLE OF INVENTION: COMPOSITIONS AND METHODS
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
                                                                                                                                    ADDRESSEE: THE SCRIPPS RESEARCH INSTITUTE STREET: 10666 No. 6074872th Torrey Pines I CITY: La Jolla STATE: California COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          8 CALALLGLALAICS-QGAASQPDLDLASRRLLQRALAAA
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Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 554-2937
TELEFAX: (619) 554-6312
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 110 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acids

TOPOLOGY: linear

MOLECULE TYPE: protein

FRAGMENT TYPE: C-terminal

US-08-648-322-3
                                                                                                                                                       STRL
CITY: La
STATE: Calif
COUNTRY: US
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Length 2476;

DB 2; .6e+02; es 24;

Gaps

28;

Length 396;

DB 4;

Gaps

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30 LDLASRRLLQRALAAALPHRSGVSERWRTFYPNCPCLRWRPRKVKGPQLKAK
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Embarcadero Center
                                                                                                                                                                                                                                                                                                                                                   10;
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NUMBER: 36,627
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION
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                                                                           TELECOMMUNICATION
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US-08-484-105-10
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                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                   RESULT 8

US-09-632-098-2
; Sequence 2, Application US/09632098
; Patent No. 6420154
; GENERAL INFORMATION:
; APPLICANT: Sheppard, Paul O.
; APPLICANT: Baindur, Nand
; APPLICANT: Bishop, Paul D.
; TITLE OF INVENTION: MAMMALIAN ADHESION PROTEASE PEPTIDES
; FILE REFERENCE: 99-39
; CURRENT APPLICATION NUMBER: US/09/632,098
; CURRENT FILING DATE: 2000-08-02
; NUMBER OF SEQ ID NOS: 26
; SOFTWARE: FastSEQ for Windows Version 3:0
; SEQ ID NO 2
; LENGTH: 802
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-632-098-2
11 CCLALLALPAQSCGPGRGPVGRRRYARKQLVPLLYKQFVPGVPERTLG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            th 12.9%; Score 58; DB 4; Length Similarity 30.0%; Pred. No. 58; 24; Conservative 5; Mismatches 25; Ind
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: ORIGIN OF REPLICATION COMPLEX GE
3: 24
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ZIP: 94111-4187.
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/484,105
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US-08-484-105-10
; Sequence 10, Application US/08484105
; Patent No. 5589341
: GENERAL INFORMATION:
APPLICANT: STILLMAN, Bruce
APPLICANT: KOBAYASHI, RYUJI
APPLICANT: RINE, Jasper
; APPLICANT: ROSS, Margit
; APPLICANT: FOSS, Margit
; APPLICANT: LAURENSON, Patricia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TEST,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MCNALLY, Francis J
LAURENSON, Patricia
HERSKOWITZ, Ira-
LI, Joachim J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: FLEHR, HOHBACH, TES
STREET: 4 Embarcadero Center,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ,
10,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            an Francisco
California
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APPLICANT: RINE, Jase
APPLICANT: FOSS, Marg
APPLICANT: MCNALLY, E
APPLICANT: LAURENSON,
APPLICANT: LAURENSON,
APPLICANT: LI, Joachi
APPLICANT: CI, Joachi
APPLICANT: CI, Kin
TITLE OF INVENTION: C
                               SERWRTFYPN 62
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71 SERFRDLVPN 80
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Best Local S
Matches 24
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                       ABBOTT PARK
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Best Local Similarity
Matches 16; Conser
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US-08-469-260A-344
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                                                                                                                      COMPUTER:
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                                                        COUNTRY
                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION
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                                                                                                                                                                                                       USOUT 11
USOUT 129

SEQUENCE 29, Application US/09143571

Patent No. 6333153

GENERAL INFORMATION:

APPLICANT: FISHEL, Richard

APPLICANT: GRADIA, Scott

APPLICANT: GRADIA, Scott

TILE OF INVENTION: COMPOSITIONS, KITS, AND METHODS FOR EFFEC

TILE OF INVENTION: WCLEOTIDE MODULATION OF DNA MISMATCH REC

TILE OF INVENTION: PROTEINS

FILE REFERENCE: 9855-601

CURRENT APPLICATION NUMBER: US/09/143,571

CURRENT APPLICATION NUMBER: 60/066,977

EARLIER FILING DATE: 1997-01

SARLIER APPLICATION NUMBER: 60/066,977

EARLIER FILING DATE: 1997-08-28

NUMBER OF SEQ ID NOS: 36

SOFTWARE: PATENTIN VOY: 2.0

SECO ID NOS: 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: JOHN N. SIMONS
APPLICANT: TAMI J. PILOT-MATIAS
APPLICANT: TAMI J. PILOT-MATIAS
APPLICANT: GEORGE J. DAWSON
APPLICANT: GEORGE G. SCHLAUDER
APPLICANT: SURESH M. DESAI
APPLICANT: THOMAS P. LEARY
APPLICANT: ANTHONY SCOTT MUERHOFF
APPLICANT: JAMES C. ERKER
APPLICANT: SHERI L. BUIJK
APPLICANT: SHERI L. BUIJK
APPLICANT: ISA K. MUSHAHWAR
TITLE OF INVENTION: NON-A, NON-B. NON-D, NON-E HEPPI
TITLE OF INVENTION: REAGENTS AND METHODS FOR THEIR USE
NUMBER OF SEQUENCES: 716
                                                                                                                                    ---VQYKLKTLYPNIPTTDYDPLQVEEPFLL
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                                                  Score
Pred.
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CORRESPONDENCE ADDRESS:
ADDRESSEE: ABBOTT LABORATORIES
                                                                               10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 25.7%; Pu
Matches 28; Conservative 12;
                                                 12:7%;
30.8%;
                                                                Similarity 30.8 16; Conservative
    protein
                                                                                                                                65 VELVSWKPLLQAIART--
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-143-571-29
/ MOLECULE TYPE:
US-08-484-106-10 ;
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US-08-469-260A-344
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ENGTH: 834
                                                Query.Match
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Matches 16
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Length 47;
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                                         APPLICATION DATA:
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30-DEC-1993
                                                                                                       US/08/424,550
                               C-DOS/MS-DOS
Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                            Application US/08176427B
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                    compatible
PC-DOS/MS
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IM: PC-DOS/MS
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Proteins
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                                                                                                                                                                      ATION INFORMATION 708-937-6365 708-938-2623
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            disk
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laritý 41.0%;
Conservative
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Tabin, Cliffor
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FORM:
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and Uses Related Thereto
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14-DEC-1994
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                                                                                                                                                       Application US/08460900C
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llarity 28.6%;
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                                                                                                                                                      th 396;
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APPLICANT: McMahon, Andrew P.
APPLICANT: Tabin, Clifford J.
TITLE OF INVENTION: Vertebrate Embryonic Pattern-Inducir TITLE OF INVENTION: Proteins and Uses Related Thereto NUMBER OF SEQUENCES: 47
CORRESPONDENCE ADDRESS:
ADDRESSEE: LAHIVE & COCKFIELD
STREET: 60 State Street
                                                                                                                                                                                   Indels
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Mismatches
                                                                                                                                                  th 12.6%; Score 56.5; I Similarity 28.6%; Pred. No. 37; 20; Conservative 7; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: BOS.L...
STATE: MA
COUNTRY: USA
ZIP: 02109
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: ASCII(text)
CURRENT APPLICATION.DATA:
APPLICATION NUMBER: US/08/356,060A
FILING DATE: 14-DEC-1994
CLASSIFICATION 435
PRIOR APPLICATION NUMBER: US 08/176,427
FILING DATE: 30-DEC-1993
ATTORNEY/AGENT INFORMATION:
NAME: Vincent, Matthew P.
REGISTRATION NUMBER: 36,709
REFERENCE/DOCKET NUMBER: HMI-006CP
TELEPHONE: (617) 227-7400
TELEPHONE: (617) 227-7400
TELEPHONE: (617) 227-5941
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARRACTERISTICS:
LENGTH: 396 amino acids
TYPE: amino acid
TYPE: amino acid
TYPE: amino acid
TYPE: ALINE PROCEETIFE: PROCEETI
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Pred. No. 37
                                                                                                                                                                                                              8 CALALLGLALAICSOGAASQPDLDLASRRLL---
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                                                                                                                                                                                                                                                                                                                                                                              Sequence 9, Application US/08356060A Patent No. 5844079 GENERAL INFORMATION: APPLICANT: Ingham, Phillip W.
                                                                                                                                                                               7;
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Best Local Similarity 28.6%;
Matches 20; Conservative 7
TELEPHONE: (617) 227-7400
TELEFAX: (617) 227-5941
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
                                                             396 amino acids
                                                                                                         protein
                                                                                                                                                                                                                                                                      53 SERWRTFYPN 62
||||:|| ||
71 SERFRDLVPN 80 ·
                                                                                          linear
                                                                                        TOPOLOGY: II
                                                                            amino
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S-08-356-060A-9
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Best Local S
Matches 20
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Gaps

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Search completed: March 21, 2003, 11:42:33 Job time : 20.2403 secs

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450
1 MRVSQIHCALALLGLALAIC.....RWRPRKVKGPQLKAKEDLER
                                                                                              March 21, 2003, 11:35:23; Search time 53.1502 (without alignments) 215.607 Million cell
GenCore version 5.1.4_p5_4578
(c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                                                                                                                                                                               Total number of hits satisfying chosen parameters:
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                                                                    protein search, using sw model
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Gapop 10.0 , Gapext 0.5
                 Copyright
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Sequence:
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Title

8470 90 Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries Minimum DB seq length: 0 Maximum DB seq length: 2000000000

A_Geneseq_101002:*

| SIDS1/gcgdata/geneseq/geneseqp-emb1/AA1980.
| SIDS1/gcgdata/geneseq/geneseqp-emb1/AA1981.
| SIDS1/gcgdata/geneseq/geneseqp-emb1/AA1983.
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| SIDS1/gcgdata/geneseq/geneseqp-emb1/AA1986.
| SIDS1/gcgdata/geneseq/geneseqp-emb1/AA1986.
| SIDS1/gcgdata/geneseq/geneseqp-emb1/AA1986.
| SIDS1/gcgdata/geneseq/geneseqp-emb1/AA1986.
| SIDS1/gcgdata/geneseq/geneseqp-emb1/AA1998.
| SIDS1/gcgdata/geneseq/geneseqp-emb1/AA1999.
| SIDS1/gcgdata/geneseq/geneseqp-emb1/AA1991.
| SIDS1/gcgdata/geneseq/geneseqp-emb1/AA1991.
| SIDS1/gcgdata/geneseq/geneseqp-emb1/AA1996.
| SIDS1/gcgdata/geneseq/geneseqp-emb1/AA1998.
| SIDS1/gcgdata/geneseq/geneseqp-emb1/AA19998.
| SIDS1/gcgdata/geneseq/geneseqp-emb1/AA1998.
| SIDS1/gcgdata/geneseq/geneseqp-emb1/AA1998.
| SIDS1/gcgdata/geneseq/geneseqp-emb1/AA1998.
| SIDS1/gcgdata/geneseq/geneseqp-emb1/AA 'AA2002

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being print and is derived by analysis of the total score distribution.

SUMMARIES

	Description	Rainbow trout prep	Rainbow trout prep	Rainbow trout prep	Human secreted pro	Novel human diagno	Angiotensin conver	Sequence of prepro	Propionibacterium	Novel human secret	Peptide #10276 enc
						• •.					
	ID	AAU07668	AAU07667	AAU07666	AAG03774	ABG08924	AAU02976	AAP20029	AAU55408	AAU30069	ABB42770
	DB	22	22	22	21	22	22	m	22	22	22
	ry ch Length DB	111	. 115	114	116	124	517	125	190	814	95
ф	Query	100.0	78.6	19.0	17:7	15.7	15.6	15.4	15.3	15.3	15.0
	Score	450	353.5	85.5	79.5	0	70	69.5	69	69	67.5
	Result No.		7	m	4	ις.	9	7	α	თ	10

pro-sequence"

"PPSS-II''

note

"Mature PPSS-II''"

note-

"Prosomatostatin II'

note

Cleavage-site

Peptide

Peptide

	1n #8062	Human brain expres	Human bone marrow	341	Peptide #10620 enc	n peptide e	bacteriu	Human MDDT SEQ ID	nitro	Bovine endothelial	Human secreted pro	poly	_	ņ	ene 29 e	ecret	Human ORFX protein	Novel human diagno	enyl	roxyphen	ostatin-2	Propionibacterium	isseria gonorrh	Murine MSH5 (mMSH5	bact	Human ORFX protein	Human PRO4400 poly	04400	~Н	Human polypeptide,	E	Herbicidally activ	Human RinTinTin pr	C	Novel human diagno	
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	11	12	13	14	. 15	16	17	. 18.	19	20	21	. 22	23	24	. 25	26	27	. 28	. 29	. 30	. 31	32	33	34	35	. 36	37	38	39	40	41	. 42	43	44	4.5	

ALIGNMENTS

RESULT 1

pPSS-I; coma; AIDS; mellitus; disorder; HIV; hypersecretion (PPSS-II'') polypeptide anticonvuls ioot; ease "PPSS-II'' pre-sequence" ..25 note=."Signal peptide" trout preprosomatostatin II Location/Qualifiers etastasis; gene immunodeficiency entry) gastroenteropancreatic note (first mykiss standard; Oncorhynchus 04-DEC-2001 neoplasm; r anti-human AAU07668; AAU07668 Key Peptide Rainbow Rainbow Protein Protein .rag

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in polypeptide
or a portion of
their
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               yonists are effective in the encoding the polypeptides are can be targeted to neoplasms of their secretory products.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       s and as adjuncts growth hormone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    agonist useful
                                 glucagonoma; AIDS diabetes mellitus;
                                                                                                                                                                        ;; nootropic; cytostatic;
anticonvulsant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cause tumour shrinkage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to an Oncorhynchus mykiss somatostatin polypepti containing a portion of preprosomatostatin I (PPSS-II). The protein sequences and their associated polynucleotides are useful for identifying modified somatostatin polypeptide which functions as a somatostatin agonist use for research, therapeutics or diagnostics, including medical and veterinary applications. The wild-type somatostatin and its modified version are useful for treating hypersecretion from endocrine tumours the pituitary (e.g. acromegaly) or gastroenteropancreatic tissues (e.g gastrinoma, glucagonoma, carcinoid syndrome), to cause tumour shrinkag through their effects on cell proliferation and apoptosis and as adjunin the treatment of diabetes mellitus via inhibition of growth hormone
                                                                                                                       disorder;
                                                                                                                                              neuroprotective
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  II' protein.
specifically claimed in the
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acquired
       hypersecretion;
gland; glucaqonor
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                                                                                                                       neurological
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New somatostatin polypeptides derived from Oncorhynchus for treating diabetes mellitus, acromegaly, gastrinoma, immunodeficiency syndrome and neurological disorders -
                                                            y; gastrinoma;
apoptosis; gro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                somatostatin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (e.g. epilepsy, Alzheimer's and somatostatin antagonists
                                                                                                                                            Alzheimer's disease; Huntington's diseas
metastasis; gene therapy; antidiabetic;
immunodeficiency virus; osteopathic; ar
   atin; preprosomatostatin;
ocrine tumour; pituitary (
tissue; acromegaly; gastr
1 proliferation; apoptosis
                                                                                                                                                                                                     osteopathic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           can
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e= "PPSS-II' pro-sequence"
                                                                                                                   syndrome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "Prosomatostatin II'"
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ts O. Mykiss PPSS-II'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               peptides
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                                                                                                                                                                                                                                                                                                            Location/Qualifiers
1..25
/note= "Signal peptide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "Mature PPSS-II'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   115
- "SS-14 variant
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                                                                                                                 1mmunodeficiency
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/ith acquired
                 PPSS-II'; PPSS-II'; enacedastroenteropancreatic to the stroenteropancreatic to the strong term of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                 700te="1
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This sequence represents
Note: The features for t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Moore CA,
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                                                                                                                                                                                                                                                                                                                                                                                       1..87
/note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            88..10
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                                                                                                                                                                                                                                                            mykiss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2001-425997/46.
B; AAS12934.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Misc-difference
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                                                                                                                                                                                                                                                            Oncorhynchus
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                                                                                                                                                                                                   antī-human
                                                                                                                 glucagon;
epilepsy;
                                                                                                                                                                     neoplasm;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5;
                                                                                                                                                                                                                                                                                                              Key
Peptide
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N-PSDB;
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Note: The features for this sequence are specifically claimed in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   agonist useful
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acquired
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100.0%; Score 450; DB 22; Lengt
Best Local Similarity 100.0%; Pred. No. 3.2e-49;
Matches 86; Conservative 0; Mismatches 0; Indel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New somatostatin polypeptides derived from Oncorhynchus for treating diabetes mellitus, acromegaly, gastrinoma, immunodeficiency syndrome and neurological disorders -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rainbow trout preprosomatostatin II (PPSS-II') polypept
"Dibasic cleavage site"
   /note= "Dibasic cleavage site"
98.111
/note= "SS-14 variant peptide"
                                                                                                                                                                                                                                                                                                                                                                                                Kittelson JD;
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AAU07667 standard; Protein; 115 AA.
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N-PSDB; AAS12935.
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for research, therapeutics or diagnostics, including medical and veterinary applications. The wild-type somatostatin and its modified version are useful for treating hypersecretion from endocrine tumours in the pituitary (e.g. acromegaly) or gastroenteropancreatic tissues (e.g. gastrinoma, glucagonoma, carcinoid syndrome), to cause tumour shrinkage through their effects on cell proliferation and apoptosis and as adjuncts in the treatment of diabetes mellitus via inhibition of growth hormone
                                                                              as a somatostatin agonist useful
                                                                                                                                                                                                                                                                                                                                                                  secretory products
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 of a large needing. The 5'
                                                                                                                                                                                                                                                                                     disease and are effective
                                                                                                                                                                                                               treatment of diabetes mellitus via innibition. Juragion secretion is addition, dysfunctional somatostatin secretion ated with acquired immunodeficiency syndrome (AIDS) and valued with acquired immunodeficiency syndrome (AIDS) and valued with acquired immunodeficiency syndrome (AIDS) and valued in antagonists are effective aton's disease) and somatostatin antagonists are effective
                                                           modified
                                                                                                                                                                                                                                                                                                                                                                                                          claimed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 114;
                                        equences and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sequence is a polypeptide encoded by one lerived from mRNAs encoding secreted prote from total human RNAs or polyA+ RNAs d
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.0065;
13;
                                                                                                                                                                                                                                                                                                                                            gene therapy and fusion peptides can be metastases, inhibiting the release of tence represents O. Mykiss PPSS-I protein
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                                                                                                                                                                                                                                                                                                                           leic acids
peptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 85.5; D Pred. No. 0.005; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        expressed sequence chromosome mapping.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                mapping
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                                        preprosomatostatin associated polynucleotides are somatostatin polypeptide which for research, therapeutics or consearch, the will be consearch.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 13; SEQ ID 7855; 71pp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       th 19.0%;
Similarity 52.3%;
23; Conservative
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                                                                                                                                                                                                                                                                                  neurological disorder
Huntington's disease)
treatment of such con
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-500381/45
N-PSDB; AAC03780.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               . 114 AA;
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Note: The featur
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clated with
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their
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Matches
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                                                                               Gaps
                                                                                                              Length 115;
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                                                                             6; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Rainbow trout; somatostatin; preprosomatostatin; hype PPSS-II'; PPSS-II'; endocrine tumour; pituitary gland gastroenteropancreatic tissue; acromegaly; gastrinoma carcinoid syndrome; cell proliferation; apoptosis; grapucagon; acquired immunodeficiency syndrome; neuroloepilepsy; Alzheimer's disease; Huntington's disease; neoplasm; metastasis; gene therapy; antidiabetic; noo anti-human immunodeficiency virus; osteopathic; antic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New somatostatin polypeptides derived from Oncorhynch for treating diabetes mellitus, acromegaly, gastrinom immunodeficiency syndrome and neurological disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Rainbow trout preprosomatostatin I (PPSS-I) polypepti
                                         22;
                                        Score 353.5; DB
Pred. No. 6e-37;
; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          pre-sequence"
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.te= "PPSS-I pro-sequence"
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/note= "Prosomatostatin I"
99.100
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/note= "Dibasic cleavage
101..114
/note= "SS-14 peptide"
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e= "Mature PPSS-I"
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                                                                                                                                                                                                                                                                                                                                         AAU07666 standard; Protein; 114 AA
                                                                                                                                                                                                                       61 PNCPCL--RPRKVKCP-AGAKEDL 81
                                                                                                                                                                                                    PNCPCLRWRPRKVKGPQLKAKEDL 84
                                                                    5.
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                                  78.6%;
illarity 86.9%;
Conservative
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89..100
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89
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N-PSDB; AAS12933.
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                                       cal Similarity 73; Consociation
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Sequence
                                     Query Match
Best Local
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Peptide
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mutations
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                                                                                                                                                                                                            116;
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different tissues. EST sequences usually correspond mai untranslated region (UTR) of the mRNA because they are from oligo-dT primed cDNA libraries. Such ESTs are not isolating cDNA sequences derived from the 5' ends of mR those cases where longer cDNA sequences have been obtainthose cases where longer cDNA sequences have been obtainthose cases where longer cDNA sequences have been obtain UTR is rarely included. 5' ESTs are derived from mRNAs ends and can therefore be used to obtain full length cD DNAs. 5' ESTs are also used in diagnostic, forensic, ge chromosome mapping procedures. They are used to obtain regulatory sequences and to design expression and secre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to isolated polynucleotide (I) and polypeptide (II) sequences. (I) is useful as hybridisationly polymerase chain reaction (PCR) primers, oligomers, and and gene mapping, and in recombinant production of (II) polynucleotides are also used in diagnostics as expresse for identifying expressed genes. (I) is useful in gene to restore normal activity of (II) or to treat disease (II). (II) is useful for generating antibodies against guantitating a polypeptide in tissue, as molecular weigh
                                                                                                                                                                                                      Query Match
Best Local Similarity 34.9%; Pred. No. 0.038;
Matches. 29; Conservative 13; Mismatches 28; Indel
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         mapping; gene mapping; gene therapy medical imaging; diagnostic; genetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 20; SEQ ID No 39283; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel human diagnostic protein #8915.
                                                                                                                                                                                                                                                                                                                                                                                                                                              ABG08924 standard; Protein; 124 AA.
                                                                                                                                                                                                                                                                                                                            64 PCLRWRPRKVKGPQLKAKEDLER 86
| | | : | | : | | | :
56 --LLSEPNQTENDALE-PEDLSO 75
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2000US-0649167
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N-PSDB; AAS73111.
                                                                                                                                                                           116 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; chromosome supplement; m
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23-AUG-2000;
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cardiovascular disease;
medical
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leep vein thrombosis;
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                                                                                                  invention.
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                                                                                                                                                                                                                  43;
                                                                      products
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                        expression
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Mismatches
                                                                        and to produce other types of data and pamino acid sequences. ABG00010-ABG30377 diagnostic amino acid sequences of the Note: The sequence data for this patent specification, but was obtained in elect
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 disease;
                  ide and polynucleotide seque forensics, gene mapping, i for genetic disorders or of ce other turn.
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                                                                                                                        specification, but was obtained in
at ftp.wipo.int/pub/published_pct
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                                                                                                                                                                                     th 15.7%;
Similarity 31.0%;
26; Conservative
                                                                                                                                                                                                                                                                                                                                                                                    AAU02976 standard; Protein;
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99IL-0133455
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multiple sclerosis; in
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                    The polypeptide and diagnostics, forential
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N-PSDB; AAS06076.
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capable of binding
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                                                     ulagnostics, tol
responsible for
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supple of s
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10-DEC-1999;
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Matches
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AAU55408
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    -LP
                                                                                                                                                                                                                                                                                                                                                                                          18;
The sequence represents an angiotensin converting enzyme spl (ACEV) polypeptide. The polypeptides of the invention includ granulocyte colony stimulating factor receptor, glucagon, in platelet-derived endothelial cell growth factor, cyclin-depending inhibitor IC, cellular tumour antigen P53, and vasoactive in polypeptide receptor 2. The polypeptides and their associate acids are useful for identification of variant sequences and candidate compounds capable of binding the molecules. The set the invention can be used in the treatment and diagnosis of disorders including cardiovascular diseases such as arterios myocardial infarction and coronary arterial thrombosis, rena such as diabetic nephropathy, muscular diseases such as hype immune disorders such as immune complex nephritis, multiple cancer, sarcoidosis, nonarcoidotic pulmonary granulomatous das asbestosis and vascular pathologies involving an endothel
                                                                                                                                                                                                                                                                                                                                                                                                                                                     :| :|
SFRLRHP
                                                                                                                                                                                                                                                                                                                                                    ore 70; DB 22; Length 517; ed. No. 3.7; Mismatches 21; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The inventors claim preprosomatostatin-1, prosomatostatin-1
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112.125
/label Somatostatir
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Best Local Similarity 37.8%; Pred.
Matches 28; Conservative 7; Mis
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    W-RPRKVKGPQLKA 80
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AAN20034.
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14-AUG-1992
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N-PSDB;
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Protein
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1017 represent Proplonibacterium acnes immunogenic ins and their associated DNA sequences are used in an and diagnosis of medical conditions caused by include SAPHO syndrome (synovitis, acne, and osteomyelitis), uveitis and endophthalmitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        osteomyelitis;
ystem; ELISA;
sorbent assay;
                                                                                    125 AA
                                                                                                                                                                                                                                                                            Gaps
  and DNA
                                                                                                                      carboxy
                                                psequently cleaved to yield S1 prepro S1 is probably about 20 predicts the sequence of a 125 is a 14 AA sequence at its carbox only 2 AAs, and is termed
                                                                                                                                                                                                                                                                                                          -PDLDLASRR--LLQRALAAALPHRSGVSER
                                                                                                                                                                                                                                                                                                                                              -LSOE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         c acids useful fo especially useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ral nervous system; Ilinked immunosorbent
                                                                                                                                                                                                                                                                          15;
                                                                                                                                                                                                                                          Length 125;
 prosomatostatin-2 and somatostatin-2; canslation of somatostatin mRNA yields containing a signal peptide which may cansit into the endoplasmic reticulum,
                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      pustulosis; hypertosis;
joint; central nervous s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       protein #16304
                                                                                                                                                                      field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       s and nucleic
infections, es
                                                                                                                                                                                                                                                                        17;
                                                                                                                                                                                                                                         3,
                                                                il) is subsequently ortion of prepro Si plas2 predicts the contains a 14 AA s
                                                                                                                                                                                                                                        Score 69.5; DB Pred. No. 0.78; ; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ston; acne vulgaris; enzyme l
osteopathic; neuroprotectant
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Carter
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                                                                                                                                        S1 by. only
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diagnosing in
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mitcham
                                                           resultant precursor (pro S1) is siteslf. The prepeptide portion obases long. Translation of pLaS2 peptide which surprisingly contaiterminus which differs from S1 by Somatostatin 2 (S2).

(Updated on 16-AUG-2002 to add mi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        acne;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Jen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  standard; Protein; 190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        pone;
                                                                                                                                                                                                                                                                      7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequences AAU39105-AAU68017 repropriets. The proteins and the treatment, prevention and continuates. The disorders included.
                                                                                                                                                                                                                                         15.4%;
36.1%;
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                                                the transit
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2000US-208841P
2000US-216747P
                                                                                                                                                                                                                                                                                                          IHCALALLGLALAICSOGAASO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SAPHO syndrome; synovitis; uveitis; endophthalmitis; tinflammatory lesion; acne v
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                 Query Match
Best Local Similarity 36.1
Matches 22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Persing DH,
e J, Zhang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DH,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Propionibacterium acnes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQ ID No
preprosomatostatin-2
encoding them. The
precursor (prepro S1
released during the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-616774/71
N-PSDB; AAS59569.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Propionibacterium
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          dermatological;
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02-JUN-2000;
07-JUL-2000;
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pustulosis,
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e central
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P. acnes is also involved in infections of bone, joints and nervous system, however it is particularly involved in the lilesions associated with acne vulgaris. A method for detecting presence or absence of P. acnes in a patient comprises contagample with a binding agent that binds to the proteins of the and determining the amount of bound protein in the sample. The polypeptides may be used as antigens in the production of an specific for P. acnes proteins. These antibodies can be used downregulate expression and activity of P. acnes polypeptides therefore treat P. acnes infections. The antibodies may also diagnostic agents for determining P. acnes presence, for example and inked immunosorbent assay (ELISA).

Note: The sequence data for this patent did not form part of specification, but was obtained in electronic format directly at ftp.wipo.int/pub/published_pot_sequences.
                                                                                                                                                                                                                                                                                                                         190;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to novel human secreted polypeptide polypeptides are usefudelypeptides and antibodies to the polypeptides are usefudetermining the presence of or predisposition to a diseas with altered levels of polypeptide. The polypeptides are identifying agents (agonists and antagonists) that bind the expressing the proteins are useful for identifying a there
                                                                                                                                                                                                                                                                                                                                                                                          11 ALLGLALAICSOG--AASQPDLDLASRRLLQRALAAAL------P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; vaccination; gene therapy; nutritional supplemenstem cell proliferation; haematopoiesis; nerve tissue rimmune suppression; immune stimulation; anti-inflammato
                                                                                                                                                                                                                                                                                                                                                          22; Indel
                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 25.8%; Pred. No. 1.5;
Matches 25; Conservative 10; Mismatches 22; Indel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        || | |
|-----APCRRRDTRQRIVVWHLPVASHYASWRPRVVR 124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ---WRPRKVK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel human secreted protein #560.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18-APR-2000; 2000US-0552929. 26-JAN-2001; 2001US-0770160.
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                                                                                                                                                                                                                                                                                    190 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 58 TFYPNCPCLR--
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expression or
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predicting
                                                                                regeneration;
                                                                                 sue growth or regeneration anti-inflammatory agents; 3304 represent the amino a s of the invention.
                                                                     and
                                                                                                                                                                                                                                        SELHIILGLFGLLLLTCLCGTAWLCCAPTGRIPSGOVSQTQLTAA - - WAPGCPQSWRRM
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                      genetically
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                                        vaccination, test
lonal supplements.
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                      and
                      peptides and
useful for
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          tide.
                                                                                  ssue
                                                                                                                                                                                8.6;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 acid probes nexpression in
                                                                                                                                                                  Score 69; DB Pred. No. 8.6; Mismatches
                                                                                                                                                                    OB
                                                                            mone, cartilage, tendon and/or nerve tissummune suppression and/or stimulation; as an treatment of leukaemias AAU29510-AAU33 equences of novel human secreted proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ome-derived single exon nucleic
gene expression in human fetal
                                                       nutritional
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                                                              l proliferation;
tendon and/or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2000US-0180312.
2000US-0207456.
2000US-0608408.
2000US-0632366.
2000US-0234687.
2000US-0234687.
                                                                                                                                                                 15.3%;
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                                                                                                                                                                                         Conservative
                                              useful
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analyzing gene expres
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                                                        can
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21-SEP-2000;
27-SEP-2000;
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ABB4277
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04-FEB-2000;
26-MAY-2000;
30-JUN-2000;
03-AUG-2000;
21-SEP-2000;
27-SEP-2000;
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the probes
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                  Query Match
Best Local S
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Matches 23
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                                                                                                                                                                                                                                                                                                                                                                                                                     Human;
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from WIPO
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                                                                                     Length
                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to single exon nucleic aci measuring human gene expression in a sample derived from ABA21535-ABA41305). The present sequence is a protein en probe. The probes may be used for predicting, measuring gene expression in samples derived from the human heart By measuring gene expression, the probes are useful for diagnosing, grading, staging, monitoring and prognosing human heart and vascular system e.g. cardiovascular dise hypertension, cardiac arrhythmias and congenital heart disconting the sequence data for this patent did not form par specification, but was obtained in electronic format dir at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     cell
  electronic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; gene expression; heart; microarray; vascular scardiovascular disease; hypertension; cardiac arrhyth congenital heart disease.
printed specification, but was obtained in electronic from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                             ----CSOGAASOPDLDLASRRLLQRAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Protein #8062 encoded by probe for measuring heart
                                                                                  th 15.0%; Score 67.5; DB 22; Similarity 27.7%; Pred. No. 1; 23; Conservative 11; Mismatches 20;
                                                                                                                                                                           Claim 15; SEQ ID No 27833; 530pp; English.
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26-MAY-2000; 2000US-0207456.
30-JUN-2000; 2000US-0608408.
03-AUG-2000; 2000US-0632366.
21-SEP-2000; 2000US-0234687.
27-SEP-2000; 2000US-0236359.
04-OCT-2000; 2000GB-0024263.
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Best Local S
Matches 23
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Pred. No. 1;
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such as Alzheimer's disease,
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microarray; Alzu
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The present sec
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specification, but was obtained in electroni
at ftp.wipo.int/pub/published_pct_sequences
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Pred. No. 1;
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expression in
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Best Local Similarity 27.7%;
Matches 23; Conservative
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analyzing gene expres
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Pred. No: 1;
11; Mismatches
                                                                                                                                                                                                                                                                                                           ton; gene (lymphoma;
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     --RIFYPNCPCLRWRPRK
                         SWSSWWDRSWWTRC---AWRPRR
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; 2000US-0207456.
; 2000US-0608408.
; 2000US-0632366.
; 2000US-0234687.
; 2000US-0236359.
; 2000GB-0024263.
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                                                                                                                                            standard; Protein;
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Similarity 27.7%;
23; Conservative
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                                                                                                                                                                                                                         (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-488900/53.
                                                                                                                                                                                                                                                                                                                           microarray; cancer;
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30-JUN-2000;
03-AUG-2000;
21-SEP-2000;
27-SEP-2000;
04-OCT-2000;
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The present invention relates to single exon nucleic see AAI31315-AAI57546). The present sequence is a peps such probe. The probes are useful for producing a mich predicting, measuring and displaying gene expression from human placenta. The probes are useful for antenathuman genetic disorders.
                                                                                                                                                                                                                                                    probes
                                                                                                                                                                                                                                                  Human genome-derived single exon nucleic acid analyzing gene expression in human placenta -
                                                                                                                                                                                                                                                                                   SEQ ID No 36852; 654pp; English.
                                                                                                                                                                                                       Rank DR;
                                                                                                                                                                                (MOLE-) MOLECULAR DYNAMICS INC.
                                                                                                                                                                                                       Chen W,
                                                                                     2000US-0180312.
2000US-0207456.
2000US-0608408.
2000US-0632366.
2000US-0234687.
2000US-0236359.
                                                                 2001WO-US00663
                                                                                                                                                                                                     Hanzel DK,
                                                                                                                                                                                                                            WPI; 2001-488897/53
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                    WO200157272-A2
                                                                                     04-FEB-2000;
26-MAY-2000;
30-JUN-2000;
03-AUG-2000;
21-SEP-2000;
27-SEP-2000;
04-OCT-2000;
Homo sapiens
                                                                 30-JAN-2001;
                                           09-AUG-2001
                                                                                                                                                                                                                                                                                    Claim 27;
                                                                                                                                                                                                                                                                                                                                                                                       Sequence
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21, 2003, 11:38:00 Search completed: March Job time : 55.1502 secs

Length 95;

DB 22;

; Score 67.5; DI ; Pred. No. 1; 11; Mismatches

Ouery Match
Best Local Similarity 27.7%;
Matches 23; Conservative

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Indels

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55 RW----RTFYPNCPCLRWRPRK 72

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-739b-18 rapb

us-09-727

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GenCore version 5.1.4_p5_4578

Copyright (c) 1993 - 2003 Compugen Ltd.

OM protein - protein search, using sw model

Run on: March 21, 2003, 11:53:33; Search time 2.12446 Seconds (without alignments)

276.816 Million cell updates/; perfect score: 58
Sequence: 1 SVDNLPPRERK 11
Scoring table: BLOSUM62
Gapop 10.0, Gapext 0.5
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Total number of hits satisfying chosen parameters: Minimum DB seq length: 0 Maximum DB seq length: 200000000

221153 seqs, 53462247 residues

Searched:

221153

Post-processing: Minimum Match 0% Maximum.Match 100% Listing first 45 summaries Published_Applications_AA:*

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2: /cgn2_6/ptodata/1/pubpaa/PCT_NEW_PUB.pep:*
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9: /cgn2_6/ptodata/1/pubpaa/USO9_NEW_PUB.pep:*
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14: /cgn2_6/ptodata/1/pubpaa/USO0_NEW_PUB.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed and is derived by analysis of the total score distribution.

SUMMARIES

										•											
	Description	Sequence 46, Appl	343,	43933	Sequence 156, App	158,	30, Ap	104,	103,	e 1400	Sequence 47, Appl	5614	Sequence 10205, A	Sequence 20, Appl	L)	Sequence 86, Appl	Sequence 22, Appl	4	35,	6537,	
	ID	US-09-050-516-46	US-09-764-860-343	US-09-864-761-43933	US-10-062-254-156	US-10-062-254-158	US-09-291-417-30	US-09-533-029-104	US-09-291-417-103	US-09-815-242-14001	US-09-050-516-47	US-09-815-242-5614.	US-09-815-242-10205	US-09-828-523A-20	US-09-815-242-12527.	US-09-828-523A-86	US-09-792-200B-22	US-09-050-516-42	US-09-916-790-35	US-09-738-626-6537	•
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APPLICATION DATA:

PRIOR

APPLICATION NUMBER

Sequence 1, Appli	Sequence 5, Appli	8	4869,	10	4, Ap	e^{-1198}	122,	e 122,	122,	41,	174,	e 174,	247,	e 247,	e 35, Ap	e 1356	e. 54,	97,	146,	Sequence 6025, Ap	e 3, 2	2, Ap	47,	Sequence 8, Appli	46
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-12	-115-8	-868-	-09-815-242-486	US-09-815-242-10798	US-10-036-729-4	US-09-815-242-11982	10-025-380-12	-09-922-217	US-09-833-263-122	-866-582-4	1-48	S-09-764-846	1-48	S-09-764-846-2	٠	-09-815-2	5-09-768-826-5	-542-2	-012-542-	-626-	US-09-802-741A-3	US-09-431-226-2	US-09-909-567B-47	8-831	US-09-864-761-46557
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ALIGNMENT

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T APPLICATION DATA:
ICATION NUMBER: US/09/050,516
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cation US/09050516
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                                     ING-MEDEL,
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MEDIUM TYPE: Diskette
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SVDNLPPRERK 11
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FEATURE:
NAME/KEY: SITE
LOCATION: (53)
OTHER INFORMATION: Xaa equals any of the naturally occurring
NAME/KEY: SITE
LOCATION: (97)
OTHER INFORMATION: Xaa equals any of the naturally occurring
1S-09-764-860-343
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 2
US-09-764-860-343
is Sequence 343, Application US/09764860
js Patent No. US20020094953A1
js GENERAL INFORMATION:
js APPLICANT: Rosen et al.
jt TILE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
js FILE REFERENCE: PC008
js CURRENT APPLICATION NUMBER: US/09/764,860
js CURRENT FILING DATE: 2001-01-17
js Prior application data removed - consult PALM or file wrapper
js SOFTWARE: Patentin Ver. 2.0
js SEQ ID NO 343
js LENGTH: 197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 3.
US-09-864-761-43933
; Sequence 43933, Application US/09864761
; Patent No. US20020048763A1
; GENERAL INFORMATION:
; APPLICANT: Penn, Sharron G.
; APPLICANT: Hanzel, David K.
; APPLICANT: Chen, Wensheng
; TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC
; TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
; FILE REFERENCE: Aeomica-X<sup>1</sup>1
; CURRENT APPLICATION NUMBER: US/09/864,761
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                                                                                                                                                                                                                                                                                                                                                        ore 38; DB 10; ed. No. 2.3; ... Mismatches
                                                                                                                        TELEX:
TELEX:
INFORMATION FOR SEQ ID NO: 46:
SEQUENCE CHARACTERISTICS:
LENGTH: 33 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: No. US20010010904Ale
US-09-050-516-46
                                                                                                                                                                                                                                                                                                                                                    Ouery Match 65.5%; Score 38; Best Local Similarity 63.6%; Pred. No. Matches 7; Conservative 2; Mismatch
NAME: Becker, Cheryl L:
REGISTRATION NUMBER: 35,441
REFERENCE/DOCKET NUMBER: 606:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/935-1729
TELEFAX: 847/938-2623
                                                                                                                                                                                                                                                                                                                                                                                                                                          1 SVDNLPPRERK 11
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N: PCT/US01/00662 01-30

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                                                                                                          Sequence Listing Engine
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01-30
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Pred. No.
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                                                 09/608,408
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Best Local Similarity 54.
Matches 6; Conservative
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FEATURE:
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Polynucleotides Encoding Proteins Involved In Plant Metaboli
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NVENTION: STE20-RELATED PROTEIN KINASES
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ER: 60/3
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APPLICANT: L1, Zhongsen
APPLICANT: Miao, Guo-Hua
APPLICANT: Miao, Guo-Hua
APPLICANT: Niu, Xiping
APPLICANT: Niu, Xiping
APPLICANT: Old. Joan
APPLICANT: Sakal, Hajime
APPLICANT: Sakal, Hajime
APPLICANT: Sakal, Hajime
APPLICANT: Sakal, Hajime
APPLICANT: Shu, Qun
TITLE OF INVENTION: POLYNUCLEOLIDES Encoding Proteins Inv
FILE REFERENCE: 2002-02-01
PRIOR PREPILATION NUMBER: US/10/062,254
CURRENT FILING DATE: 2000-07-28
PRIOR APPLICATION NUMBER: 60/146511
PRIOR FILING DATE: 1999-07-30
PRIOR FILING DATE: 1999-09-30
PRIOR FILING DATE: 1999-09-30
PRIOR FILING DATE: 1999-09-30
PRIOR FILING DATE: 1999-10-01
PRIOR FILING DATE: 1999-10-01
PRIOR FILING DATE: 1999-12-09
PRIOR FILING DATE: 1999-12-09
PRIOR FILING DATE: 1999-12-09
PRIOR FILING DATE: 1999-12-22
PRIOR FILING DATE: 1999-12-23
PRIOR FILING DATE: 1999-12-30
PRIOR PR
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Pred. No. 65;
; Mismatches 1; Ind
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US-10-062-254-158
Sequence 158, Application US/10062254
Patent No. US20020138882A1
GENERAL INFORMATION:
APPLICANT: Cahoon, Edgar B
APPLICANT: Falco, Saverio Carl
APPLICANT: Fang, Yiwen
APPLICANT: Hantke, Sabine S.
APPLICANT: Lee, Jian-Ming
APPLICANT: Lee, Jian-Ming
APPLICANT: Miao, Guo-Hua
APPLICANT: Miao, Guo-Hua
APPLICANT: Morgante, Michele
APPLICANT: Niu, Xiping
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Odell, Joan
Rafalski, Antoni
Sakai, Hajime
Zheng, Peizhong
Zhu, Qun
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SLDHLPPSEQ 18
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; NAME/KEY: UNSURE
; LOCATION: (107)
US-10-062-254-156
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Identification of Essential Genes in Prokaryotes
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                            , Application US/09815242
0020061569A1
                                                                                                                                                                             2000-03-21
2000-03-21
NUMBER: 60/191,078
                                                                                                                                                                                                                                                                                                                                                   Version
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US20010010904A1
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                                                                            Zyskind, Judith W
Wall, Daniel
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KRATOCHVIL, JO
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Best Local Similarity 85.
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                                                                                                    Frawick, John
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                                                                                                                                                                                              TON NUMBER
                                                                                                            Grant
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US-09-815-242-14001
                                                                                                                         Yamamoto,
                                                             selbeck,
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|DSLPPRE 748
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               Sequence 14001
Patent No
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US-09-050-516-47
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RESULT 8
US-09-291-417-103
US-09-291-417-103
Sequence 103, Application US/09291417A
Publication No. US20030050230A1
GENERAL INFORMATION:
APPLICANT: MARTINEZ, RICARDO
APPLICANT: WHYTE, DAVID
TITLE OF INVENTION: STE20-RELATED PROTEIN KINASES
FILE REFERENCE: 240/300
CURRENT APPLICATION NUMBER: US/09/291,417A
CURRENT FILING DATE: 1999-04-13
EARLIER FILING DATE: 1998-04-14
NUMBER OF SEQ ID NOS: 147
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 103
LENGTH: 591
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 60.3%; Score 35; DB 9; Lot 100.0%; Pred. No. 2.1e+02; ive 0; Mismatches 0;
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Pred. No. 1.4e+02;
1; Mismatches 0
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                                                                                                                                                                                                                                                                                               ORGANISM: Arabidopsis thaliana FEATURE:
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Best Local Similarity 85.7%;
Matches 6; Conservative
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6; Conservative
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S-09-533-029-104
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US/09/815,242
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FastSEQ for Windows Version
10205
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                                                                                                                                                 ; ORGANISM: Staphylococcus aureus US-09-815-242-5614
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-242-10205
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US-09-815-242-5614
Sequence 5614, Application US/09815242
Sequence 5614, Application US/09815242
Patent No. US20020061569A1
GENERAL INFORMATION:
APPLICANT: Haselbeck, Robert
APPLICANT: Oliver, John D.
APPLICANT: Trawick, John D.
APPLICANT: Yamamoto, Robert T.
APPLICANT: Yamamoto, Robert T.
APPLICANT: Au, H. Howard
TITLE OF INVENTION: Identification of Essential Genes in TITLE OF INVENTION: Prokaryotes
FILE REFERENCE: ELITRA.011A
CURRENT APPLICATION NUMBER: US/09/815,242
CURRENT APPLICATION NUMBER: 60/191,078
PRIOR FILING DATE: 2000-03-21
PRIOR FILING DATE: 2000-03-21
PRIOR FILING DATE: 2000-03-21
PRIOR FILING DATE: 2000-05-23
                                           STATE: 12
COUNTRY: USA
ZIP: 60064-3500
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 34; DB 10;
Pred. No. 99;
2; Mismatches
                                                                                                                                                                                                                                                                 CLASSIFICATION:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/828,855
FILING DATE: 31-MAR-1997
ATTORNEY/AGENT INFORMATION:
NAME: Becker, Cheryl L.
REGISTRATION NUMBER: 35,441
REFERENCE/DOCKET NUMBER: 6065.US.P1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/935-1729
TELEFAX: 847/938-2623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     No. US20010010904Ale
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R APPLICATION NUMBER: 60/242,578

R FILING DATE: 2000-10-23

R APPLICATION NUMBER: 60/253,625

R FILING DATE: 2000-11-27

R APPLICATION NUMBER: 60/257,931

R FILING DATE: 2000-12-22
STREET: 100 Abbott Park Road CITY: Abbott Park STATE: 1L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: 47
SEQUENCE CHARACTERISTICS:
LENGTH: 215 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: NO. US200100
US-09-050-516-47
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Best Local Similarity 60.0%;
Matches 6; Conservative
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Length 299;

harmacia & Upjohn Company ON: ANTIMICROBIAL METHODS AND MATERIALS 268.62120101

Pharmacia

US/09828523A

NUMBER: US/09/828,523A : 2001-04-06

aureus coding region cloned

sequence encoded by in E. coli.

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DB 9; L. 1.5e+02; 0;

Score 34; DB Pred. No. 1.5 0; Mismatches

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us-09-727-739b-18.rapb

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OTHER INFORMATION: Amino acid
OTHER INFORMATION: expression
-09-828-523A-86
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Best Local Similarity 100.0%;
Matches 6; Conservative
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SOFTWARE: Pat
SEQ ID NO 86
LENGTH: 311
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APPLICANT: Xu, H. Howard
TITLE OF INVENTION: Identification of Essential Genes in
TITLE OF INVENTION: Prokaryotes
FILE REFERENCE: ELITRA.011A
CURRENT APPLICATION NUMBER: US/09/815,242
CURRENT FILING DATE: 2001-03-21
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Pred. No. 1.5e+02;
Mismatches 0;
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Pred. No. 1.5e+02;
Mismatches 0;
APPLICANT: The Pharmacia & Upjohn Company.

TITLE OF INVENTION: ANTIMICROBIAL METHODS AND
FILE REFERENCE: 268.62120101
CURRENT APPLICATION NUMBER: US/09/828,523A
CURRENT FILING DATE: 2001-04-06
PRIOR APPLICATION NUMBER: 60/266,327
PRIOR FILING DATE: 2000-04-06
NUMBER OF SEQ ID NOS: 99
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 60/191,078
PRIOR FILING DATE: 2001-03-21
PRIOR FILING DATE: 2000-03-21
PRIOR FILING DATE: 2000-05-23
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR FILING DATE: 2000-10-23
PRIOR FILING DATE: 2000-10-23
PRIOR FILING DATE: 2000-10-23
PRIOR FILING DATE: 2000-11-27
PRIOR APPLICATION NUMBER: 60/253,625
PRIOR FILING DATE: 2000-11-27
PRIOR FILING DATE: 2000-12-22
PRIOR FILING DATE: 2001-02-16
NUMBER OF SEQ ID NOS: 14110
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 12527
LENGTH: 303
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3-09-815-242-12527
Sequence 12527, Application US/09815242
Patent No. US20020061569A1
PRIENT INFORMATION:
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100.0%; Pi
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US-09-815-242-12527
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T: Ohlsen, Kari L.
T: Zyskind, Judith W.
T: Wall, Daniel
T: Trawick, John D.
T: Carr, Grant J.
T: Yamamoto, Robert T.
                                                                                                                                                                                                                                                                                                       Similarity 100
6; Conservative
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39 VDNLPP
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LENGTH: 303
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O'KEEFE, DANIEL
VENTION: SOYBEAN GLUTATHIONE-S-TRANSFERASE ENZYMES
NCE: CL-1108-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 219;
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US-08-344-536-2

US-08-344-536-2

US-08-920-562-2

US-08-920-562-2

US-08-920-562-2

US-08-976-626A-3

US-09-147-915-4

US-09-028-934-30

US-08-624-663A-2

US-08-974-565C-1
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                                                                                                                                                                                                                                                                 ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANI:
TITLE OF INVENTION: SOYBEAN GLUIA....
FILE REFERENCE: CL-1108-A
CURRENT APPLICATION NUMBER: US/09/247,373B
CURRENT FILING DATE: 1999-02-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 39;
Pred. No.
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1999-02-10
ER: 08/924.747
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Microsoft Office
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PRIOR APPLICATION NUMBER:
PRIOR FILING DATE: 1997.
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NUMBER OF SEQ ID NOS:
SOFTWARE: Microsoft
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TITLE OF INVENTION:
FILE REFERENCE: CL
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DNLPPRDK
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SOFTWARE: M1
SEQ ID NO 52
LENGTH: 219
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SEQ ID NO 34
LENGTH: 224
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LOW TEMPERATURE-SENSITIVE FERMENTABILI
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Pred. No.
                                                                                                                 ; ORGANISM: Staphylococcus epidermidis US-09-134-001C-3917
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Plaza
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          , ORGANISM: Saccharomyes cerevisiae
US-09-125-028-2
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I, Yasuhiro
Kozo
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ilarity 66.7%;
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30 Rockefeller
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US-09-134-001C-3917
Sequence 3917, Application US/09134001C
Patent No. 6380370
GENERAL INFORMATION:
APPLICANT: Lynn Doucette-Stamm et al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING
TITLE OF INVENTION: EPIDERMIDIS FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: GTC-007
CURRENT APPLICATION NUMBER: US/09/134,001C
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US-09-134-001C-5085
US-09-134-001C-5085
Sequence 5085, Application US/09134001C
Patent No. 6380370
GENERAL INFORMATION:
TITLE OF INVENTION:
TITLE OF INVENTION: EPIDERMIDIS FOR DIAGNOSTICS AND THERAPED
FILE REFERENCE: GTC-007
CURRENT APPLICATION NUMBER: US/09/134,001C
CURRENT APPLICATION NUMBER: US 60/064,964
PRIOR APPLICATION NUMBER: US 60/064,964
PRIOR FILING DATE: 1997-11-08
PRIOR FILING DATE: 1997-11-08
PRIOR FILING DATE: 1997-08-14
NUMBER OF SEQ ID NOS: 5674
SEQ ID NO 5085
LENGTH: 248
LENGTH: 248
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5169835-6
; Patent No. 5169835
; TITLE OF INVENTION: PREGNANCY SPECIFIC PROTEINS APPLICAT
; NUMBER OF SEQUENCES: 48
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/07/390,409
; FILING DATE: 07-AUG-1989
; SEQ ID NO.66:
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 Length 2
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Pred. No. 11;
1; Mismatches
 Score 39; DB 4
Pred. No. 10;
2; Mismatches
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Pred. No. 45;
4; Mismatches
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Similarity 75.0%; P. 6; Conservative 2;
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Best Local Similarity 70.0%;
Matches 7; Conservative
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Compositions
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Release #1.0,
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PAK4; A No
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APPLICATION NUMBER: U
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ATION NUMBER:
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                                                                                                                                                                                               TITLE OF INVENTION:
TITLE OF INVENTION:
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New York
                   Similarity
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GENERAL INFORMATION:
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US-09-082-737-2
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5169835
; Patent No. 5169835
; APPLICANT: WAI-YEE, CHAN
; TITLE OF INVENTION: PREGNANCY SPECIFIC PROTEINS APPLICATIONS
; NUMBER OF SEQUENCES: 48
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/07/390,409
; TIING DATE: 07-AUG-1989
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  MEDIUM TYPE: Diskette - 3.50 inch, 1440 Kb storage COMPUTER: IBM PS/V OPERATING SYSTEM: MS-DOS Ver3.30 SOFTWARE: PATENT AID Ver1.0 CURRENT APPLICATION DATA:
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Pred. No. 5.3e+02;
2; Mismatches 1
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Pred. No. 53;
3; Mismatches
                                                                APPLICATION NUMBER: US/08/894,344C
FILING DATE: 15-AUGUST-1997
PRIOR APPLICATION DATA:
                                                                                                      APPLICATION NUMBER: JP343700/95
FILING DATE: 28-DECEMBER-1995
APPLICATION NUMBER: PCT/JP96/03862
FILING DATE: 27-DECEMBER-1996
ATTORNEY/AGENT INFORMATION:
NAME: PERTY, LAWIENCE S:
REGISTRATION NUMBER: 31865
                                                                                                                                                                                                                                                                                                                    Saccharomyces cerevisiae
X2180-18
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 212-218-2100
TELEFAX: 212-218-2200
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                              2958 amino acids ino acid
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Best Local Similarity 66.7%;
Matches 6; Conservative
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Best Local Similarity 54.5%;
Matches 6; Conservative
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101 TINNENPRENK 111
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257 IDVLPPKER 265
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MEDIUM TYPE:
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FINTION: SOYBEAN GLUTATHIONE-S-TRANSFERASE
ICE: CL-1108-A
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CURRENT APPLICATION NUMBER: US 60/064,964

PRIOR APPLICATION NUMBER: US 60/065,779

PRIOR APPLICATION NUMBER: US 60/055,779
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1999-02-10
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Pred. No.
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Pred. No
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e 5632, Application US/09134001C
No. 6380370
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; ORGANISM: Staphylococcus epidermidis US-09-134-001C-5632
                                                                                                                                                                                                                                                                                                                                                                                                                                                      08/924,747
                                                                                                                                                                                                                                                                                 Patent No. 6168954
GENERAL INFORMATION:
APPLICANT: MCCONTAINS
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100.0%;
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SOFTWARE: Microsoft Office
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                                      223 amino acid
amino acid
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6; Conservative
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5; Conservative
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NUMBER OF SEQ ID NOS: 5674
SEQ ID NO 5632
LENGTH: 305
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200 DSLPPRDK 207
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GENERAL INFORMATION:
APPLICANT: George W. Kemble
TITLE OF INVENTION: A Novel VZV Gene, Mutant VZV and
TITLE OF INVENTION: Compositions
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
ADDRESSEE: Cooley Godward Castro Huddleson & Tatum
STREET: 5 Palo Alto Square
CITY: Palo Alto
STATE: California
COUNTRY: USA
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/857,534
FILING DATE:
CLASSIFICATION:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/04971
FILING DATE:
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78;
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/235,406
FILING DATE: APRIL 28, 1994
ATTORNEY/AGENT INFORMATION:
NAME: Luann Cserr
REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: AVIR-004/00WO
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-857-0663
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 34; DB
Pred. No. 78
0; Mismatches
                                                                                                                                                                                                                           NAME: Luann Cserr
REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: AVIR-004/00US
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-843-5165
TELEFAX: 415-857-0663
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APPLICATION NUMBER: 08/235,406
FILING DATE:
ATTORNEY/AGENT INFORMATION:
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223 amino acids

TOPOLOGY: unknown

, MOLECULE TYPE: Prot-

US-08-857-534-12
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Best Local Similarity 100.0%;
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 223 amino acids
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3: sp_fung1:*
4: sp_human:*
5: sp_mammal:*
7: sp_mhc:*
8: sp_phage:*
9: sp_plant:*
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2: sp_virus:*
3: sp_vortebrate:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed and is derived by analysis of the total score distribution.

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ce sturgeon).
Craniata; Vertebrata; Euteleostomi;
Acipenseriformes; Acipenseridae;
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EMBL; AF395849; AAL13248.1; -.

InterPro; IPR004250.
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D65FBD7C6F1E4E4D CRC64;
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Actinopterygii; Neopterygii; Teleostei; Osteoglosso
Osteoglossiformes; Notopteridae; Chitala.
              Osteoglossum
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somatostatin cDN
identification
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Pred. No. 0.02;
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             Osteoglossidae;
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Ctinopterygii; Chondrostei;
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larity 52.08;
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AF292650; AAK9
            Osteoglossiformes;
NCBI_TaxID=109271;
                                                                                                                  osteoglossomorphs:
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CBI_TaxID=7904;
                                                                              Al-Mahrouki A
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Pfam; PFO?
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Q90Y40
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.

Al-Mahrouki A.A., Irwin D.M., Youson J.H.;

Al-Mahrouki A.A., Irwin D.M., Youson J.H.;

Characterization of variant somatostatin cDNAs from several costeoglossomorphs: molecular identification and comparative analysis.";

Submitted (AUG-2000) to the EMBL/GenBank/DDBJ databases.

EMBL; AF292652; AAK97069.1;

EMBL; AF292652; AAK97069.1;

InterPro; IPR004250; Somatostatin.

R InterPro; IPR004250; Somatostatin; 1.

SEQUENCE 114 AA; 12494 MW; 454DA57A309CA8F2 CRC64;
01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
Preprosomatostatin.
Pantodon buchholtzi (Butterflyfish).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutelee
Actinopterygii; Neopterygii; Teleostei; Osteoglossomorpha;
Osteoglossiformes; Pantodontidae; Pantodon.
NCBI_TaxID=8276;
[1]
SEQUENCE FROM N.A.
Al-Mahrouki A.A., Irwin D.M., Youson J.H.;
Characterization of variant somatostatin cDNAs from sever;
Characterization of variant identification and comparative
osteoglossomorphs: molecular identification and comparative
analysis.";
Submitted (AUG-2000) to the EMBL/GenBank/DDBJ databases.
EMBL; AF292651; AAK97068 1;
InterPro; IPR004250; Somatostatin.
InterPro; IPR004250; Somatostatin.
Pfam; PF03002; Somatostatin; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           090Y43
090Y43;
090Y43;
01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-DEC-2002 (TrEMBLrel. 20, Last annotation update)
01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
Preprosomatostatin.
Preprosomatostatin.
Osteoglossum bicirrhosum (silver arawana).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eute
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      090Y41
090Y41
090Y41;
090Y41;
01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-DEC-2002 (TrEMBLrel. 20, Last annotation update)
01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
Preprosomatostatin.
Gnathonemus petersii.
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutelinopterygii; Neopterygii; Teleostei; Osteoglossomorplosteoglossiformes; Mormyridae; Gnathonemus.
NCBI_TaxID-42645;
                                                                                                                                                                                                                                                                                                                                  Ouery Match
Best Local Similarity 52.0%; Pred. No. 10.00095;
Matches 13; Conservative 5; Mismatches 7; Indel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indel
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Best Local Similarity 64.0%; Pred. No. 0.02;
Matches 16; Conservative 2; Mismatches 7
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| |: |:| || || || || || :|
1 MKLCQVHCILALLGLVLGMCGSSSA 25
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Q90Y43
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Sollars C., Vallarino M

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116. AA

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Indels

·Length 116;

13;

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Euteleostomi

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Osteoglossomorpha;

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MAGNESIUM IS REQUIRED FOR CATALYSIS AND FOR STABILIZING (BY SIMILARITY).
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                                                                                      PHOSPHOENOLPYRUVATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 132,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                O'Meara B.C., Normark B.B.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           agriculture in beetles (Curculionidae:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                               (enolase)
                                 ., Sonthayanon B.;
hydratase (enolase
                                                                                                                                                                                                                                                                                                                 Pyruvate.
EB575C6FB541ABD3 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CRC64;
                                                                        Pac. J. Mol. Biol. Biotechnol. 7:89-94(1999)
CATALYTIC ACTIVITY: 2-PHOSPHO-D-GLYCERATE - P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BCB814053FFA3471
                                                                                                                                                          MODIMER (BY SIMILARITY).
LOCATION: CYTOPLASMIC (BY SIM BELONGS TO THE ENOLASE FAMILY AAC78141.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ore 52; DB ed. No. 3.6; Mismatches
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Last annotation
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S
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laga; Scolytidae;
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                                             phosphopyruvate
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Pred.
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lase; 1.
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                                                                                                                                                                                                                                                                                                                Lyase; Magnesium
434 AA; 47265 MW
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                                                                                                                                                                                                                                 Enolase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Enolase; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 42.6%;
                                                                                                                                                         SUBCELLULAR LOCATION: C
                                                                                                                                                                                                                                                                         Enolase;
                                                                                                                                                                                                                                                                                                  S00164; ENOLASE;
                                                                                                                                                                                                                                                                                                                                                                                                                                 108 ANAILGVSLAICKAGAA 124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Phytophaga;
                                                                                                                                                                                                                                                                                                                                                                        Similarity 64.
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                                                                                                                                       PATHWAY: GLYCOLYSIS.
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                                                          rawn, Penaeus
c. J. Mol. Bio
                                                                                                                                                                                                                              IPR000941;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Neoptera;
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                                 Boons
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10; Conser
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ICBI_TaxID=141164;
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SEQUENCE
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HSSP; P5
InterPro
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SEQUENCE FROM N.A.
Al-Mahrouki A.A., Irwin D.M., Youson J.H.;
"Characterization of variant somatostatin cDNAs from several costeoglossomorphs: molecular identification and comparative analysis.";
Submitted (AUG-2000) to the EMBL/GenBank/DDBJ databases.
EMBL; AF292653; AAK97070.1;
InterPro; IPR004250; Somatostatin.
Pfam; PF03002; Somatostatin; 1.
SEQUENCE 114 AA; 12561 MW; 4E3C32F58E34F971 CRC64;
                                                                                                                                                           h 114
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Q96316

Q96316,

Q96316,

Q96316,

Q1-DEC-2001 (TrEMBLrel. 19, Created)

O1-DEC-2001 (TrEMBLrel. 19, Last sequence update)

O1-DEC-2001 (TrEMBLrel. 20, Last annotation update)

I 01-MAR-2002 (TrEMBLrel. 20, Last annotation update)

Enolase (Fragment).

Aphanarthrum sp. SCR03.

Bukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Inse

Envolution

S Eukaryota; Meoptera; Endopterygota; Coleoptera; Polyphaga

Cucujiformia; Phytophaga; Scolytidae; Aphanarthrum.

NCBI_TaxID=113463;

NCBI_TaxID=113463;

I 1]

SEQUENCE FROM N.A.

Farrell B.D., Sequeira A.S., O'Meara B.C., Normark B.B.,

Jordal B.H.;

and Platypodinae).";

EMBL; AF375332; AAK54790.1;

EMBL; AF375332; AAK54790.1;

Pram; PF00113; enolase; 1.

Probom; PD000902; Enolase; 1.

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                                                                                                                                             uery Match
est Local Similarity 56.0%; Pred. No. 0.82;
atches 14; Conservative 3; Mismatches 8; Indels
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O96656;
O96656;
O1-MAY-1999 (TrEMBLrel. 10, Created)
O1-MAY-1999 (TrEMBLrel. 10, Last sequence update)
O1-JUN-2002 (TrEMBLrel. 21, Last annotation update)
O1-JUN-2002 (TrEMBLrel. 21, Last annotation update)
Enolase (EC 4.2.1.11) (2-phosphoglycerate dehydratase) (3 glycerate hydro-lyase).
Penaeus monodon (Penoeid shrimp).
Eukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca; Eumalacostraca; Eucarida; Decapoda; Dendrobranchiata; Pen NCBI_TaxID-6687;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 54; DB 5; Length
Pred. No. 2.8;
4; Mismatches 3; Inde
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22700 MW; 34E3408775FED864 CRC64
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| ::| |||||| ||| : | ||
| Db. 1 MLSTRIQCALALLSLALPVSSVYAA 25
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Best Local Similarity 58.8%;
Matches 10; Conservative
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| |:||::||:| |||
79 ATAILGVSLAVCKAGAA 95
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SEQUENCE 216 AA;
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096656
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Dendroctonus mexicanus.
Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;
Pterygota; Neoptera; Endopterygota; Coleoptera; Polyphaga;
Cucujiformia; Phytophaga; Scolytidae; Dendroctonus.
NCBI_TaxID=77163;
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Pfam; PF00113; enclara
                                                                                                                                                                                                                        (TrEMBLrel. (TrEMBLrel. (TrEMBLrel.
                                                                                                                                                                          PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE
STRAIN=HLT16;
Sequeira A.S., Farrell E
"Evolutionary origins of
heetle herbivores?";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          00113; enola PD0000902; En
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096310;
01-DEC-2001
01-DEC-2001
01-MAR-2002
                                                                                                                                                                                                                         |-DEC-2001
|-DEC-2001
|-MAR-2002
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Best Local S
Matches 10
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Pfam; PF(
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SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Scolytinae
CO963H3
PRELIMINARY;
PRT; 133 AA.
CO963H3;
CO963H3;
CO963H3;
TO1-DEC-2001 (TrEMBLrel. 19, Created)
TO1-DEC-2001 (TrEMBLrel. 19, Last sequence update)
TO1-MAR-2002 (TrEMBLrel. 20, Last annotation update)
E Enolase (Fragment).
N ENOINI.
S SCOlytus unispinosus.
S SCOlytus unispinosus.
C Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;
Pterygota; Neoptera; Endopterygota; Coleoptera; Polyphaga;
Cucujiformia; Phytophaga; Scolytidae; Scolytus.
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096313;
01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-DEC-2002 (TrEMBLrel. 21, Last annotation update)
101-JUN-2002 (TrEMBLrel. 21, Last annotation update)
11 Enolase (Fragment).
12 Enolase (Fragment).
13 Enolase (Fragment).
14 Enolase (Fragment).
15 Enolase (Fragment).
16 Enolase (Fragment).
17 Enolase (Fragment).
18 Enolase (Fragment).
19 Enolase (Fragment).
10 Enolase (Fragment).
10 Enolase (Fragment).
11 Enolase (Fragment).
12 Enolase (Fragment).
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"The evolution of agriculture in beetles (Curculionidae and Platypodinae).";

Evolution 0:0-0(2001).

EMBL; AF375336; AAK54794.1; -
InterPro; IPR000941; Enolase.

Pfam; PF00113; enolase; 1.

PRINTS; PR01610; CD36ANTIGEN.

ProDom; PD0009902; Enolase; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                             Jordal B.H.;
"The evolution of agriculture in beetles (Curculionidae and Platypodinae).";
Evolution 0:0-0(2001).

EMBL; AF375346; AAK54804 1; -.

InterPro; IPR000941; Enolase.

Pfam; PF00113; enolase; 1.

ProDom; PD000902; Enolase; 1.
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Farrell B.D., Sequeira A.S., O'Meara B.C., Normark B.B.
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Farrell B.D., Sequeira A.S., O'Meara B.C., Normark B.B.
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13879 MW; BF8D60C5D97AC008 CRC64;
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Best Local Similarity 58.8%;
Matches 10; Conservative
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28D98F57787CC168 CRC64;

15194 MW;

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B.D.; of Gondwanan interactions: How

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Enolase;

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4; Mismatches
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Pro; IPR000941; Enolase
PF00113; enolase; 1,
PF0000902; Enolase; 1,
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Similarity 58.8%;
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Length 149;

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Similarity
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O1-DEC-2001 (TrEMBLrel. 19, Created)
O1-DEC-2001 (TrEMBLrel. 19, Last sequence update)
O1-DEC-2001 (TrEMBLrel. 19, Last sequence update)
O1-MAR-2002 (TrEMBLrel. 20, Last annotation update)
F 01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
F 01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
F Enolase (Fragment).

N Hylastes porculus.
SEQUENCE:
FRAGORIA: Neoptera; Endopterygota; Coleoptera; Polyphage
Cucujiformia; Phytophaga; Scolytidae; Hylastes.
Cucujiformia; Phytophaga; Scolytidae; Hylastes.

N CBI_TAXID=141065;
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096311;
01-DEC-2001 (TrEMBLrel. 19, Created)
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01-DEC-2002 (TrEMBLrel. 20, Last annotation update)
Enolase (Fragment).
Enolus: (Fragment).
Eukaryota: Metazoa: Arthropoda; Tracheata: Hexapoda; Inserygota: Neoptera; Endopterygota: Coleoptera: Polyphag.
Cucujiformia: Phytophaga: Scolytidae: Dendroctonus.
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"The evolution of agriculture in beetles (Curculionidae: and Platypodinae).";

Evolution 0:0-0(2001;
Evolution 0:0-0(2001;
EMBL; AF375338; AAK54796.1;
InterPro; IPR000941; Enolase.
Pfam; PF00113; enolase; 1.
ProDom; PD000902; Enolase; 1.
NON_TER 1.
NON_TER 149 149
SEQUENCE 149 AA; 15570 MW; B09B142A0850D265 CRC64;
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NON_TER 148 148
SEQUENCE 148 AA; 15404 MW; 633A7D8C7B3D5990 CRC64
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| 20 ANAILGVSLAVCKAGAA 36
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Copyright (c) 1993 - 2003 Compugen Ltd.
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- protein search, using sw model OM protein

March 21, 2003, 11:35:23; Search time 3.00429 Seconds (without alignments) 345.142 Million cell updates Run on:

Title: Perfect score: Sequence:

US-09-727-739B-19 122 1 MRVSQIHCALALLGLALAICSQGAA 25

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

112892 seqs, 41476328 residues Searched:

112892 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

SwissProt_40:* Database : Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being print and is derived by analysis of the total score distribution.

SUMMARIES

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ALIGNMENTS

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		16-OCT-2001	16-OCT-2001 (Rel. 40, Last seq	sequence .update)		
	DI	16-OCT-2001	(Rel. 40, Last	otation	update)	,
		Somatostatin	n II precursor	tains:	[Tyr21,Gly24]somatostatin-28	in-28;
	DE 0	[Tyr7,Gly10]	somatostatin-14		-	•
		Oncornynchus	s mykiss (Rainbo	w trout) ((Rainbow trout) (Salmo gairdneri).	•
	ဗ ဗ	Eukaryota; Met	azoa;	a; Craniat	a; Vertebrata; Euteleos	tomi;
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	R.P.	ENCE F	ROM N.A.			
	.RX	95	92	28684;		
		Moore C.A.,	Kittilson J.D.,	Dah]	, Sheridan M.A.;	
		"Isolation a	and characterization	110	CDNA encoding for	
		preprosomate	cont	ng [Tyr7,	ĭ	from the
		endocrine pa	ř	ow trout,	Oncorhynchus mykiss.";	
		en. Comp.). Sg.	53-261(199		:
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	FT	۵۱	104. 115	BY	SIMILARITY.	
	SO	SEQUENCE 1	115 AA; 12963 MW	. 2	025FCA6D91 CRC64;	
	n O	Query Match	90.28;		0; DB 1; Length 115;	•
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	ζ'n	1 MKVSQIHCA	HCALALLGLALAICSQGAA	GAA 25		

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Hormone; Signal; Multigene family
                                  Otto C.J., Peter R.E.;
"The expression of SRIF mRNA in the brain of goldfish.";
Submitted (SEP-1997) to the EMBL/GenBank/DDBJ databases.
-!- FUNCTION: SOMATOSTATIN INHIBITS THE RELEASE OF SOMATOTROPIN.
-!- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY.
                                                                                                                                                            This SWISS-PROT entry is copyright. It is produced through a between the Swiss Institute of Bioinformatics and the EMBI the European Bioinformatics Institute. There are no restrictuse by non-profit institutions as long as its content is modified and this statement is not removed. Usage by and institutes requires a license agreement (See http://www.isb-sikor send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                 [TYR21,GLY24]SOMATOSTATIN-2:
[TYR7,GLY10]SOMATOSTATIN-14
BY SIMILARITY.
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98957D68011A651A CRC64;
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pancreatic
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0.0009;
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Taylor W.L., Magazin M.D., Tavi
Dixon J.E.;
ure of cloned DNA complementary
n-14 messenger RNA ".
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Pred. No.
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channel catfish.
188:19-29(1985).
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57:10372-10377(1982).
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Ictalurus.
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Similarity 54.2%;
13; Conservative
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MEDLINE-85303576; Pubme
Dixon.J.E., Andrews P.C
"Somatostatins of the c
Adv. Exp. Med. Biol. 18
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EMBL; AF025686; AAF153
InterPro; IPR004250; S
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SMS2_CARAU STANDARD; PRT; 120 AA.

O9YGH4; O9PTU2;

16-OCT-2001 (Rel. 40, Last sequence update)

16-OCT-2001 (Rel. 40, Last annotation update)

16-OCT-2001 (Rel. 40, Last annotation update)

Somatostatin II precursor [Contains: [Tyr21,Gly24]somatostatin-14].

[Tyr7,Gly10]somatostatin-14].

Carassius auratus (Goldfish).

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Actinopterygii; Neopterygii; Teleostei; Ostariophysi; Cypriniformes;

Cyprinidae; Carassius.

NCBI_TaxID=7957;
                                                                                                                                                                                                                                                                                                                                            OTROPIN
                                                                                                                                                                                                                                   SEQUENCE FACTOR SECONDARY SECONDARY PETER R.E.;

Lin X.-W., Peter R.E.;

"Cloning and characterization of CDNAs encoding preprosomatosts and -II from goldfish brain.";

L Submitted (NOV-1995) to the EMBL/GenBank/DDBJ databases.

L SUBMITTERITY INHIBITS THE RELEASE OF SOMATOTROPING OF SOMATOSTATIN INHIBITS THE RELEASE OF SOMATOTROPING OF SOMATOSTATIN FAMILY.

-I- SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY.

C This SWISS-PROT entry is copyright. It is produced through a copeween the Swiss Institute of Bioinformatics and the EMBL of the European Bioinformatics Institute. There are no restrict use by non-profit institutions as long as its content is modified and this statement is not removed. Usage by and for entities requires a license agreement (See http://www.isb-sib.ch).
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SOMATOSTATIN-26 (POTENTIAL).
SOMATOSTATIN-14.
BY SIMILARITY.
4; B5920015E2D272A4 CRC64;
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Pred. No. 0.00062;
3; Mismatches 6; Indels
                                                                              Carassius auratus (Goldfish).
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Eute
Actinopterygii; Neopterygii; Teleostei; Ostariophysi; Cy
Cyprinidae; Carassius.
NCBI_TaxID=7957;
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InterPro; IPR004250; Somatostatin.
Pfam; PF03002; Somatostatin; 1.
Cleavage on pair of basic residues; Hormone; Signal; Mul
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                      09YGH5;
16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Somatostatin IA precursor (Contains: Somatostatin-26;
  .114 AA
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Best Local Similarity 64.0%;
Matches 16; Conservative
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P SEQUENCE OF 101-114.

X MEDLINE-8126423; Pubmed-6114953;
Andrews P.C., Dixon J.E.;
Tisolation and structure of a peptide hormone predicted from a mF sequence. A second somatostatin from the catfish pancreas.";
J. Bitol. Chem. 256:1827 (1981).
C.I. BIOLICHEL LOCATION: SCHETCH INTIBILITY THE RELEASE OF SOMATOTROPIN.
C.I. SUBCELLULAR LOCATION: Secreted.
C.I. SUBCELLULAR LOCATION: Secreted.
C.I. SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY.
C.I. SIMILARITY: BELONGS TO THE SOMATOSTATIN-14.

EMBL: WOOGOB: CAA23877.1;
C.I. SEMBL: WOOGOB: CAA23877.1;
C.I. SIMILARITY: BENDABLE.
C.I. SIMILARITY: BENDABLE.
C.I. SIMILARITY: BENDABLE.
C.I. SIMILARITY: BENDABLE.
C.I. SEMBL: WOOGOB: CAA23877.1;
C.I. SIMILARITY: BENDABLE.
C.I. SIMILARITY: BENDAME.
C.I. SEQUENCE SURFACE SOMATOSTATIN-14.
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anidae; Rana
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MEDLINE-97057290; PubMed-8901629;
TOSTIVINT H., Lihrmann I., Bucharles C., Vieau D., Coulous Fournier A., Conlon J.M., Vaudry H.;
"Occurrence of two somatostatin variants in the frog brain characterization of the cDNAs, distribution of the mRNAs, receptor-binding affinities of the peptides.";
Proc. Natl. Acad. Sci. U.S.A. 93:12605-12610(1996).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
52.5%; Score 64; DB 1; Length 11
Best Local Similarity 52.0%; Pred. No. 0.017;
Matches 13; Conservative 5; Mismatches 7; Indels
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MEDLINE=93038702; PubMed=1358069;
Vaudry H., Chartrel N., Conlon J.M.;
"Isolation of [Pro2,Met13]somatostatin-14 and somatost frog brain reveals the existence of a somatostatin ger
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SMS1_RANRI STANDARD; PRT; 115 AA.
P87384; Q9PS18;
16-OCT-2001 (Rel. 40, Last sequence update)
16-OCT-2001 (Rel. 40, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Somatostatin 1 precursor (PSS1) [Contains: Somatostat.
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FEE0F2C76F74D99F CRC64
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Rana ridibunda (Laughing frog) (Marsh frog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Amphibia; Batrachia; Anura; Neobatrachia; Ranoidea;
NCBI_TaxID=8406;
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                      SOMATOTROPIN
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Segil N., Shrutkowski A., Dworkin M.B., Dworkin-Rastl E.;
"Enolase isoenzymes in adult and developing Xenopus laevis and
characterization of a cloned enolase sequence.";
Biochem. J. 251:31-39(1988).
                                                                                                                                                                                                                                                                                                                           Multigene family;
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349756FEB4ABE213 CRC64;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; l
Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea;
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  un. 188:477-482(1992).
INHIBITS THE RELEASE OF
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                                                           TO THE SOMATOSTATIN FAMILY
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SIMILARITY: BELONGS TO THE ENOLASE FAMILY.
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4.2.1.11) (2-phosphoglycerate deh
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them. Biophys. Res. Co
FUNCTION: SOMATOSTATI
SUBCELLULAR LOCATION:
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14; Conservative
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SUBUNIT: HOMODIMER.
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NCBI_TaxID=8355;
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MEDLINE=99310339; PubMed=10382966;

Kawarabayasi Y., Hino Y., Horikawa H., Yamazaki S., Haikawa Y.,

A Jin-no K., Takahashi M., Sekine M., Baba S.-I., Ankai A., Kosug
Hosoyama A., Fukui S., Nagai Y., Nishijima K., Nakazawa H.,

Takamiya M., Masuda S., Funahashi T., Tanaka T., Kudoh Y.,

Yamazaki J., Kushida N., Oguchi A., Aoki K.-I., Kubota K.,

Nakamura Y., Nomura N., Sako Y., Kikuchi H.;

"Complete genome sequence of an aerobic hyper-thermophilic

Trenarchaeon, Aeropyrum pernix Kl.";

Crenarchaeon, Aeropyrum pernix Kl.";

L. COFACTOR: TWO HEME GROUPS AND COPPER B (BY SIMILARITY).

C. I. SUBCELLULAR LOCATION: Integral membrane protein.

C. I. SIMILARITY: IN THE N-TERMINAL SECTION; BELONGS TO THE HEME

C. I. SIMILARITY: IN THE C-TERMINAL SECTION; BELONGS TO THE CYTC

C. SIMILARITY: IN THE C-TERMINAL SECTION; BELONGS TO THE EMBL

C. SIMILARITY: IN THE C-TERMINAL SECTION; BELONGS TO THE EMBL

C. SIMILARITY: IN THE C-TERMINAL SECTION; BELONGS TO THE EMBL

C. SIMILARITY: IN THE E-TERMINAL SECTION; BELONGS TO THE EMBL

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O9YDX6;
30-MAY-2000 (Rel. 39, Created)
30-MAY-2001 (Rel. 40, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Heme-copper oxidase subunit I+III (EC 1.9.3.-).
AOXB OR APE0793.
Acropyrum pernix.
Archaea; Crenarchaeota; Thermoprotei; Desulfurococcales;
Desulfurococcaceae; Aeropyrum.
NCBI_TaxID=56636;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Emthe European Bioinformatics Institute. There are no restruce by non-profit institutions as long as its content modified and this statement is not removed. Usage by and entities requires a license agreement (See http://www.isb-sor send an email to license@isb-sib.ch).
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Wakagi T., Ishikawa R.;
"Heme-copper-oxidase.";
Submitted (NOV-1998) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                       BY SIMILARITY.
MAGNESIUM (BY SIMILARITY)
MAGNESIUM (BY SIMILARITY)
MAGNESIUM (BY SIMILARITY)
83C922489B530EB4 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 54; DB 1;
Pred. No. 1.4;
; Mismatches
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PIR; S00463; NOXL.
HSSP; P56252; 1PDZ.
InterPro; IPR000941; Enolase.
Pfam; PF00113; enolase; 1.
PRINTS; PR00148; ENOLASE.
ProDom; PD000902; Enolase; 1.
TIGRFAMS; TIGR01060; eno; 1.
PROSITE; PS00164; ENOLASE; 1.
Lyase; Glycolysis; Magnesium.
INIT_MET 0 0 0
ACT_SITE 157 BY
METAL 244 244 MAN
METAL 292 292 MAN
METAL 317 317 MAN
SEQUENCE 433 AA; 47373 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Match 44.3%;
Local Similarity 64.7%;
nes 11; Conservative
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Gaps
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Galliformes; Phasianidae; Phasianinae
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                                                                                                 chain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EMBL/GenBank/DDBJ databases.
INHIBITS THE RELEASE OF SOMATOTROPIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        S., Yamamoto H.
                                                                                                  Respiratory
                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 81
                                                                                                                                                                                           (PROBABLE)
                                                                                                                                          (PROBABLE)
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Last annotation update)
c [Contains: Somatostatin-28;
                                                                                                                                                   (PRÓBABLE)
(PROBABLE)
                                                                                                                                                                       PROBABLE)
                                                                                                 Transmembrane;
                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 54; DB Pred. No. 2.4; 4; Mismatches
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IRON (F
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HSSP; P18401; 1FFT.

InterPro; 1PR000883; COX1.

InterPro; 1PR000298; CytC_oxdse_III

Pfam; PF00115; COX1; 1.

Pfam; PF00510; COX3; 1.

PRINTS; PR01165; CYCCXIDASEI.

ProDom; PD0000382; CytC_oxdse_III; 1

PROSITE; PS00077; COX1; 1.

PROSITE; PS50253; COX3; 1.

Oxidoreductase; Heme; Copper; Transr
                          oxdse_III
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Eukaryota, Metazoa; Chordata; (Archosauria; Aves; Neognathae; Gallus.
NCBI_TaxID=9031;
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larity 52.2%;
Conservative
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                                                                                                                     Somatostatin precursor
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Yonekura H., Okamoto H
Submitted (JUN-1991) t
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27,
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Best Local Similarity
Matches 12; Conser
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P33094;
01-OCT-1993
01-OCT-1993
16-OCT-2001
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COFACTOR: MAGNESIUM IS REQUIRED FOR CATALYSIS AND FOR THE DIMER (BY SIMILARITY).
PATHWAY: Glycolysis.
SUBUNIT: HOMODIMER (BY SIMILARITY).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SUBUNIT: HOMODIMER (BY SIMILARITY).
SUBCELLULAR LOCATION: CYtoplasmic.
SIMILARITY: BELONGS TO THE ENOLASE FAMILY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MAGNESIUM
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etween the Swiss Institute of Bio
he European Bioinformatics Institu
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Molecular evidence for the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Z; Enolase;
1060; er-
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ENOLASE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       94195794;
True Mar 25 08:25:48 2003

True Mar 25 08:25:40  

True Mar 25 08:25:40  

True Mar 25 08:25:48  

True Mar 25 08:
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leghorn;
5305: Pub
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PS00164; EN
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     TISSUE-Embryon
MEDLINE-890797
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5-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ProDom; PD
TIGRFAMs;
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       15-JUN-2002 (Rel. 41, Created)
15-JUN-2002 (Rel. 41, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Alpha enolase (EC 4.2.1.11) (2-phospho-D-glycerate hydro-lyase)
(Phosphopyruvate hydratase).
Alligator mississippiensis (American alligator).
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Archosauria; Crocodylidae; Alligatorinae; Alligator.
                                                                                                                                                                             TISSUE=Muscle;

MEDLINE=99439677; PubMed=10508547;

Mannen H., Li S.S.-L.;

Mannen H., Li S.S.-L.;

Mannen H., Li S.S.-L.;

Molecular evidence for a clade of turtles.";

Mol. Phylogenet. Evol. 13:144-148(1999).

-!- CATALYTIC ACTIVITY: 2-phospho-D-glycerate = phosphoenolpyr

H(2)0.

-!- COFACTOR: MAGNESIUM IS REQUIRED FOR CATALYSIS AND FOR STAB

THE DIMER (BY SIMILARITY).

-!- SUBUNIT: HOMODIMER (BY SIMILARITY).

-!- SUBCELLULAR LOCATION: Cytoplasmic.

-!- SUBCELLULAR LOCATION: Cytoplasmic.

-!- SIMILARITY: BELONGS TO THE ENOLASE FAMILY.

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01-NOV-1990 (Rel. 16, Created)
01-NOV-1990 (Rel. 16, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Alpha enolase (EC 4.2.1.11) (2-phospho-D-glycerate hydrocrystallin).
Anas platyrhynchos (Domestic duck).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eute Archosauria; Aves; Neognathae; Anseriformes; Anatidae; Al
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            n; Multigene family.
BY SIMILARITY.
MAGNESIUM (BY SIMILARITY).
MAGNESIUM (BY SIMILARITY).
MAGNESIUM (BY SIMILARITY).
A; CD3DD59FB1EBB3DA CRC64;
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Pred. No. 2.7;
4; Mismatches
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HSSP; P56252; 1PDZ.
InterPro; IPR000941; Enolase.
Pfam; PF00113; enolase; 2.
PRINTS; PR00148; ENOLASE.
TIGREAMS; TIGR01060; eno; 1.
PROSITE; PS00164; ENOLASE; 1.
Lyase; Glycolysis; Magnesium; Mul
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                47190 MW;
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| |:||::||:|
107 ANAILGVSLAVCKAGAA 123
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292
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433 AA;
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P19140;
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SEQUENCE
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REQUIRED FOR CATALYSIS AND FOR STABILIZING
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es; Phasianidae; Phasianinae;
                                          ZYME AND A LENS STRUCTURAL PROTEIN.
2-phospho-D-glycerate - phosphoenolpyruvate
and
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(2-phospho-D-glycerate hydro-lyase)
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                                                                                                                                                                                          copyright. It is produced tute of Bioinformatics and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eye lens protein
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Galliformes;
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MAGNESIUM (
                                                                                                                                                Cytoplasmic.
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244 M
292 M
317 M
47108 MW;
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Magnesium;
                                          AN ENZYME
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European Bioinformatics
                                                                                     COFACTOR: MAGNESIUM IS
                                                                                                                                             SUBCELLULAR LOCATION: C
SIMILARITY: BELONGS TO
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                                                                                                                                                                                                                                                                                                                                                                                                                  Enolase;
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             structural protein
ell Biol. 107:2729-2
FUNCTION: BOTH AN EN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STANDARD;
                                                                                                               PATHWAY: Glycolysis
SUBUNIT: HOMODIMER.
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Eukaryota, Metazoa, Cho
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-!- COFÁCTOR: MAGNESIUM IS REQUIRED FOR CATALYSIS AND FOR STABILIZING
THE DIMER (BY SIMILARITY).
-!- PATHWAY: Glycolysis.
-!- SUBUNIT: HOMODIMER (BY SIMILARITY).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15-JUN-2002 (Ref. 41, Created)
15-JUN-2002 (Rel. 41, Last sequence update)
15-JUN-2002 (Rel. 41, Last sequence update)
Alpha enclase (EC 4.2.1.11) (2-phospho-D-glycerate hydro-lyase)
(Phosphopyruvate hydratase).
Sceloporus undulatus (Eastern fence lizard) (Skink).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostom1; Sceloporus.
NCBI_TaxID=8520;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 and
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g as its content
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MAGNESIUM (BY SIMILARITY).
MAGNESIUM (BY SIMILARITY).
MAGNESIUM (BY SIMILARITY).
W; EOAD198562629D23 CRC64;
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                                                                                                                                                                                                                                                      This SWISS-PROT entry is copyright. It is produced between the Swiss Institute of Bioinformatics and the European Bioinformatics Institute. There are nuse by non-profit institutions as long as its comodified and this statement is not removed. Usage entities requires a license agreement (See http://wor send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This SWISS-PROT entry is copyright. It is produced between the Swiss Institute of Bioinformatics and the European Bioinformatics Institute. There are nuse by non-profit institutions as long as its c
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SEQUENCE FROM N.A.
HISSUE=Muscle;
MEDLINE=99439677; PubMed=10508547;
Mannen H., Li S.S.-L.;
"Molecular evidence for a clade of turtles.";
Mol Phylogenet. Evol. 13:144-148(1999).
-!- CATALYTIC ACTIVITY: 2-phospho-D-glycerate
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SUBCELLULAR LOCATION: Cytoplasmic.
SIMILARITY: BELONGS TO THE ENOLASE FAMILY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pfam; PF00113; enolase.
Pfam; PF00113; enolase; 12.
PRINTS; PR00148; ENOLASE.
ProDom; PD000902; Enolase; 1.
TIGRFAMS; TIGR01060; eno; 1.
PROSITE; PS00164; ENOLASE; 1.
Lyase; Glycolysis; Magnesium INIT_MET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    47439 MW;
                                                                                                                                                 SUBCELLULAR LOCATION: C.
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HSSP; P56252; 1PDZ.
InterPro; IPR000941; Enolase
Pfam; PF00113; enolase; 2.
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107 ANAILGVSLAVCKAGAA 123
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10; Conservative
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Best Local Similarity 42.6%; Score 52. DB 1; Length 433;
Matches 10; Conservative 4; Mismatches 3; Indels
Oy ALALGIALAICSOGAA 123

RESULT 14
BNOA_PYTRG
ID BNOA_PYTRG
ID BNOA_PYTRG
ID STANDARD; PRT; 433 AA.

C 09WTLO;
DT 15-JUN-2002 (Rel. 41, Created)
DT 15-JUN-2002 (Rel. 41, Last sequence update)
DE Alpha enolase (EC 4.2.11) (2-phospho-D-glycerate hydro-lyase)
C Python regius (Ball Python) (Royal python)
C Lepidosautia, Squamata; Scieroglossa; Serpentes; Henophidia:
C Pythonidae: Python
C C TISSIE-Munacle;
C TISSI
RA Tanaka M., Maeda K., Nakashima K.;

The Chicken alpha-enolase but not beta-enolase has a Src-deper tyrosine-phosphorylation site: cDNA cloning and nucleotide analysis."

LO Biochem. 117:554-559(1995).

CC -1 Biochem. 117:554-559(1995).

CC -1 COPACTOR: MAGNESTUM IS REQUIRED FOR CATALYSIS AND FOR THE DIMER (BY SIMILARITY).

CC -1 SUBGELLULAR LOCATION: Cytoplasmic.

-1 SIMILARITY: BELONGS TO THE ENOLASE FAMILY.

CC -1 SIMILARITY: BELONGS TO THE ENOLASE FAMILY.

-1 SIMILARITY: BELONGS TO THE ENOLASE FAMILY.

CC This SWISS-PROT entry is copyright. It is produced through the ENOPEAN BRAIN STAILCHER of Bioinformatics and the ENOPEAN BRAINFORT of Libertholms as its content of the Enopean and the statement is not removed. Usage by non-profit institutions as long as its content of modified and this statement is not removed. Usage by non-profit institutions as long as its content of modified and this statement is not removed. Usage by non-profit institutions as long as its content of modified and this statement is not removed. Usage by non-profit institutions as long as its content of modified and this statement is not removed. Usage by non-profit institutions as its content of modified and this statement is not removed. Usage by non-profit institutions as its content of modified and this statement is not removed. Usage by non-profit institutions as its content of an interproj institution of the profit institutions as its content of an interproj institution of the profit institutions as its content of an interproj institution of the profit institutions as its content of an interproj institution of the profit institution of the profit institutions as its content of the profit institutions and the profit institutions as its content of the profit institutions and the profit
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RESULT 14
ENOA_PYTRG
ID ENOA_PYTRG
ID ENOA_PY
AC 09W7L0
DT 15-JUN

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CC EMBL; AF072587; AAD41644.1; -
DR HSSP; P56252; 1PDZ.
DR InterPro; IPR000941; Enolase.
DR PRINTS; PR00113; enolase; 2.
DR PRINTS; PR00148; ENOLASE.
DR ProDom; PD000902; Enolase; 1.
DR TIGRFAMs; TIGR01060; eno; 1.
DR PROSITE; PS00164; ENOLASE; 1.
KW Lyase; Glycolysis; Magnesium; Multigene family.
FT METAL 244 MAGNESIUM (BY SIMILARITY).
FT METAL 292 292 MAGNESIUM (BY SIMILARITY).
FT METAL 317 A7362 MW; 7CD895E405529301 CRC64;
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Indel

3;

Score 52; DB 1; Pred. No. 2.7; 4; Mismatches

Query Match Best Local Similarity 58.8%; Matches 10; Conservative

9 ALALLGLALAICSOGAA 25 | |:||::||:| | 107 ANAILGVSLAVCKAGAA 123

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Search completed: March 21, 2003, 11:38:45 Job time : 5.00429 secs

Length 4

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March 21, 2003, 11:38:56; Search time 5.36481 (without alignments) 447.986 Million cell
GenCore version 5.1.4_p5_4578 Copyright (c) 1993 - 2003 Compugen Ltd.
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                                                                                                                                                                              Title: US-09-727-739B-19
Perfect score: 122
Sequence: 1 MRVSQIHCALALLGLALAICSQGAA 25
                                                                    OM protein - protein search, using sw model .
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Maximum Match 100%
Listing first 45 summaries
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Gapop 10.0 , Gapext 0.5
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Maximum DB seq length: 2000000000
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PIR_73:*
1: pirl:*
2: pir2:*
3: pir3:*
4: pir4:*

Pred. No. is the number of results predicted by chan score greater than or equal to the score of the resu and is derived by analysis of the total score distri

SUMMARIES

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		RESULT 1
		IS1064
		somatostatin ii precursor - rainbow trout C:Species: Oncorhynchus mykiss (rainbow trout)
	•	sequence_revision 1
		C; Accession: I51064 R: Moore C A : Kittilson I D : Dahl S K : Sheridan M A
		. 98, 253-261, 1995
		A; Title: Isolation and characterization of a cDNA encoding for preprosomatostatin
		A; Reference number: 151064; Muin: 95354921; FMin: 7628664 B; Accession: 151064
		A; Status: preliminary; translated from GB/EMBL/DDBJ
٠.		A; Molecule Type: mknA A: Residues: 1-115 <moo></moo>
	٠.	A; Cross-references: EMBL: U32471; NID: 9975344; PIDN: AAC59695.1; PID: 9975345
	4.	C; Superiamily: Somatostatin
		of Score
		22; Conservative 2; Mismato
		OY 1 MRVSQIHCALALLGLALAICSQGAA 25.
	:	Db 1 MKVCRIHCALALLGLALAICSOGAA 25
	• . •	
		RESULT 2
		KIIDSI somatostatin-14 precursor - channel catfish
		names: somatostatin I
		Contains: somatostatin-14
		Species: Ictalurus punctatus (channel catfish)
		C:Date: 30-Jun-1980 #sequence_revision 31-Dec-1993 #text_change 18-Jun-1999 C:Accession: S00292: A93897: A92334: A01435
		Minth, C.D.; Taylor, W.L.
		Biol. Chem. 257, 10372-10377, 1982
		A; Title: The structure of cloned DNA complementary to catfish pancreatic somatosta
		scession: S00292
		A; Molecule type: mRNA
		VIII
,		A;Cross-references: EMBL:VUUbU/; NID:gb4U1/; PIDN:CAAZ38//.1; PID:gb4U18 b.maxilor u I . Collier x I . Deschanes p I . Weith u I . Divon I F
	•	Sci. U.S.A. 78, 6694-6698, 1981
,		nalysis of a cDNA coding
	·	A93897; MUID:82082515;
		A; Accession: A93897
		A; Molecule Type: mknA A; Residues: 82-108 <tay></tay>
	-	A; Cross-references: GB: J00944

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#text_change 18-Jun-1999

clawed frog

African

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2-phospho-D-glyceric

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PIDN:CAA68706.1

glycolysis; hydro-lyase;

Asp) #status predicted

Length 434;

DB. 1;

Score Pred.

3;

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Shrutkowski, A.; Dworkin, M.B.; Dworkin-Rastl, 251, 31-39, 1988
Lase isoenzymes in adult and developing Xenopus number: S00463; MUID:88268812; PMID:3390159
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           210,343/Active site: magnesium 2 (Ser) #status predicted 245,293,318/Binding site: magnesium 1 (Asp. 210.
                                                                                                                                                                                                                                                                       laevis (African clawed frog)
#sequence_revision 31-Mar-1991
                                                                                                                                                                                                                   4.2.1.11) ENO1
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-references: EMBL:Y00718;
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ilarity 64.7%;
Conservative
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11, 1999
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A; Experimental source: strai
C; Genetics:
A; Gene: APEO793
C; Superfamily: cytochrome-c
C; Keywords: copper; electron
F; 73, 388/Binding site: heme
F; 251,300,301/Binding site:
F; 251-255/Cross-link: 1'-his
F; 255/Binding site: oxygen (
F; 386/Binding site: heme a3
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omodulator and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Species: Rana ridibunda (laughing frog)
NiAlternate names: PSS1 protein
Species: Rana ridibunda (laughing frog)
Nate: 11-Apr-1997 #sequence_revision 09-May-1997 #text_change 16-Jul-1
Nate: 11-Apr-1997 #text_change 16-Jul-1997 #t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     eprosomatostatin SS-14 - channel catfish
Species: Ictalurus punctatus (channel catfish)
C;Date: 13-Sep-1996 #sequence_revision 13-Sep-1996 #text_change
C;Accession: I50798
R;Dixon, J.E.; Andrews, P.C.
Adv. Exp. Med. Biol. 188, 19-29, 1985
A;Title: Somatostatins of the channel catfish.
A;Reference number: I50798; MUID:85303576; PMID:2863931
A;Accession: I50798
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Residues: 1-114 <DIX>
A;Residues: 1-114 <DIX>
A;Cross-references: GB:M25903; NID:g213339; PIDN:AAA49339.1; PII
C;Superfamily: somatostatin
A; Title: Isolation and structure of a peptide hormone predic A; Reference number: A92334; MUID:81264223; PMID:6114953
A; Accession: A92334
A; Molecule type: protein
A; Residues: 101-114 <AND>
C; Superfamily: somatostatin
C; Keywords: neuropeptide
F; 1-24/Domain: signal sequence #status predicted <PRO>
F; 25-100/Domain: propeptide #status predicted <PRO>
F; 25-101/Domain: propeptide #status experimental <AMATERIO3-114/Product: somatostatin-14 #status experimental
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Pred. No. 0.036;
5; Mismatches
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Pred. No. 0.27;
; Mismatches
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Local Similarity 52.0%;
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Best Local Similarity 56.0%;
Matches 14; Conservative
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Y.; Yamazaki,
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                        #text_change 20-Jun-
                                                                                                                                                                                                                                                                                               #status predicted
 - Aeropyrum
                                                            T.; Kudoh,
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                                                                                    genome sequence of an aerobic hyper-thermophilic
: A72450; MUID:99310339; PMID:10382966
                                                Haikawa
                                                                                                                                                                 PIDN:BAA79771.1;
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                                                ıki, S.;
Tanaka,
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                                                 Yamazaki
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                                                           S.; Funahashi,
 1.9.3.1)
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electron transfer; he
e: heme a iron (His)
y site: copper (His)
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a3 iron (His)
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imental source: strain Kl
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larity 52.2%;
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C; Keywords: acetyle
F; 1/Modified site:
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C;Species: Fasciola hepatica (liver fluke)
C;Date: 07-Oct-1994 #sequence_revision 07-Oct-1994 #text_change 22-Jun-1999
C;Accession: A53665
R;Davis, R.E.; Singh, H.; Botka, C.; Hardwick, C.; Ashraf el Meanawy, M.; Villanueva, J. Biol. Chem. 269, 20026-20030, 1994
A;Title: RNA trans-splicing in Fasciola hepatica. Identification of a spliced leader (A;Reference number: A53665; MUID:94327554; PMID:8051087
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               shown:
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A; Residues: 1-395 < HED>
A; Residues: 1-395 < HED>
C; Keywords: carbon-oxygen lyase; hydro-lyase
                                                                                                            Somatostatin precursor - chicken
C; Species: Gallus gallus (chicken)
C; Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_ch
C; Accession: S20630
R; Nata, K.; Kobayashi, T.; Karahashi, K.; Kato, S.; Yamamot
submitted to the EMBL Data Library, June 1991
A; Description: Nucleotide sequence determination of chicken
A; Reference number: S20630
A; Accession: S20630
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A; Molecule type: mRNA
A; Residues: 1-116 <NAT>
A; Cross-references: EMBL: X60191; NID: 962985; PIDN: CAA42747
C; Superfamily: somatostatin
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A; Residues: 1-431 <DAV>
A; Cross-references: GB:U10297; NID:9499266; PIDN:AAA57450
C; Superfamily: enolase
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5.3;
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Best Local Similarity 52.4%; Pred. No. 1.4;
Matches 11; Conservative 5; Mismatches
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Best Local Similarity 58.8
Matches 10; Conservative
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116; S38303
A.; Feo, S.
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Fitle: Complete amino acid sequence of the neurone-specific Reference number: S02077; MUID:89091176; PMID:3208766
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Ning, expression and sequence homologies of
Number: JU0060, MUID:90006764; PMID:2792767
JU0060
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/ structure of rabbit muscle enolase
A37210; MUID:91113295; PMID:2275753
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20; NID:931145; PIDN:CAA3: was confirmed by protein
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156569; S02616;
Glallongo, A.;
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Pred. No.
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Title: Complete structure of the human generate complete structure complete st
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G.; Barbieri,
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Similarity 58.8%;
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PIDN:CAA44540.1; M.; Keller, A.

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DNA cloning, PMID:2734297

CDNA

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PIDN: AAA37554.1;

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S.; Takahashi,

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PID: 9204042

PIDN: AAA41119 Suzuki, Y.

neuron-specific i, PMID:2450052

PID:9554480

hydro-lyase

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GB:M18742; NID:9205766; PIDN:AAA41725.1;
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1. Sci. U.S.A. 82, 7453-7457, 1985
Lar cloning and the nucleotide sequence of the nucleotide nu
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names: 2-phospho-glycerate dehydratase;
attus norvegicus (Norway rat)
oct-1987 #sequence_revision 25-Oct-1987
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Pred. No. 5.7;
4; Mismatches
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                                                                                     NID:950143;
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ues: 59-233,'NA',236-434 <LA2>
-references: GB:M20745; NID:g193029;
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Murine muscle-specific enolase: cl
nce number: A33921; MUID:89282789;
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site: magnesium 1
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58.88;
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A; Molecule type: DNA
A; Residues: 1-28 <SA2>
A; Residues: 1-28 
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10; Conservative
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ence number: $29675
sion: $29675
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uperfamily: enolase
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103-113,
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J. Neurosci. Res. 19, 450-456, 1988
A.Tille: Human gamma enolase: Isolation of a CDNA clone and expression in normal A. Reference number: 156599, MUID:88259288; PMID:385803
A.Status: Speliminary: translated from CB/EMBL/DDBJ
A.MOIscoule type: MRMA
A.Status: Preliminary: translated from CB/EMBL/DDBJ
A.MOIscoule type: MRMA
A.Residues: 'GC', 29-126 'N', 128-434 <VANN
A.Cross-references GB:M36768; NID:9182117; PIDN:AAA52388.1; PID:9182118
R:BBS, IR.N. A.Alsopp, M.T.B. P.; Moore, D.C.M.: Thompson, R.J.
A.Tille: Sequence conservation in the 3'-untranslated regions of neurone-specific A; Residues: 'SQ2616 'MUID:88005129; DMID:365393
A.TILLE: Sequence conservation in the 3'-untranslated regions of neurone-specific A; Residues: ACS-A34 CDAY
A.Residues: accessing MLM: Residues: ACS-A34 CDAY
A.Residues: ACS-COMEN C.R.; Oulinp, GB.; Hurt, J.; Day I.N.M.; Wischik, C.M.
Biochim: Blophys. Acta 1158, 120-128, 1993
A.Tille: Characterisation of an epitope specific to the neuron-specific isoform of blota/A4 protein
A.Residues: ACS-COMEN C.R.; Oulinp, 94 002176; PMID:7691181
A.Residues: ACS-A34 CDAY
A.Residues: ACS-A44 CDAY
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N.Alternate names: enolase beta
C; Species: Mus musculus (house mouse)
C; Date: 31-Mar-1992 #sequence_revision 31-Mar-1992 #text_change 18-C; Accession: 517109; 518036; 529675; A33921
R; Lamande, N.; Brosset, S.; Keller, A.; Lucas, M.; Lazar, M.
Submitted to the EMBL Data Library, September 1991
A; Reference number: 517109
A; Molecule type: DNA
A; Residues: 1-434 < LAM>
A; Residues: 1-434 < LAM>
A; Residues: 1-434 < LAM>
A; Cross-references: EMBL: X61600; NID: 950848; PIDN: CAA43797.1; PID: 9;
R; Peterson, C.A.; Cho, M.; Rastinejad, F.; Blau, H.M.
Submitted to the EMBL Data Library, October 1991
A; Description: Beta-enolase: a gene expressed in undifferentiated pc
A; Reference number: $18036
A; Accession: $18036
A; Molecule type: mRNA
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42.6%; Score 52; DB 1; Length 4
Best Local Similarity 58.8%; Pred. No. 5.7;
Matches 10; Conservative 4; Mismatches 3; Indel
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| 108 ANAILGVSLAVCKAGAA 124
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Phosphopyruvate hydratase (EC 4.2.1.11) alpha - duck
N;Alternate names: enolase alpha; tau-crystallin
C;Species: Anas platyrhynchos (domestic duck)
C;Date: 21-May-1990 #sequence_revision 21-May-1990 #text_change 21-Jul-2000
C;Accession: A32132
R;Wistow, G.J.; Lietman, T.; Williams, L.A.; Stapel, S.O.; de Jong, W.W.; Horwitz, J.; E
J. Cell Biol. 107, 2729-2736, 1988
A;Title: Tau-crystallin/alpha-enolase: one gene encodes both an enzyme and a lens struct
A;Reference number: A92750; MUID:89079778; PMID:2462567
A;Reference number: A92750; MUID:89079778; PMID:2462567
A;Reference number: A94 <WIS>
A;Molecule type: mRNA
A;Residues: 1-434 <WIS>
A;Residues: 1-434 <WIS>
A;Cross-references: GB:X14195; NID:962455; PIDN:CAA32409.1; PID:962456
C;Superfamily: enolase
C;Superfamily: enolase
                                                                                                                                                                                                                              lase mRNAs from mouse brain
phosphopyruvate hydratase (EC 4.2.1.11) gamma - mouse
N; Alternate names: 2-phosphoglycerate dehydratase; enolase gamma
C; Species: Mus musculus (house mouse)
C; Date: 12-Feb-1993 #sequence_revision 12-Feb-1993 #text_change 22-Jun-1999
C; Accession: S10247
R; Kaghad, M.; Dumont, X.; Chalon, P.; Lelias, J.M.; Lamande, N.; Lucas, M.; La Nucleic Acids Res. 18, 3638, 1990
A; Title: Nucleotide sequences of cDNAs alpha and gamma enolase mRNAs from mous A; Reference number: S10246; MUID: 90301487; PMID: 2362815
A; Accession: S10247
A; Molecule type: mRNA
A; Residues: 1-434 < kAGS>
A; Residues: 1-434 < kAGS>
A; Cross-references: EMBL: X52380; NID: 955494; PIDN: CAA36606.1; PID: 955495
C; Keywords: carbon-oxygen lyase; gluconeogenesis; glycolysis; hydro-lyase
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4; Mismatches
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Pred. No. 5.7;
4; Mismatches
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Job time : 7.36481 secs
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Best Local Similarity 58.8%;
Matches 10; Conservative
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2: /cgn2_6/ptodata/1/pubpaa/PCT_NEW_PUB.pep
3: /cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep
4: /cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep
5: /cgn2_6/ptodata/1/pubpaa/US07_NEW_PUB.pep
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7: /cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep
8: /cgn2_6/ptodata/1/pubpaa/US09_NEW_PUB.pep
9: /cgn2_6/ptodata/1/pubpaa/US09_NEW_PUB.pep
10: /cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep
11: /cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep
12: /cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep
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NT: LARSEN, Peter
NT: STEPHEN, John
F INVENTION: Blochemical Markers for the Human
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No. US20030044795A1
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Similarity 56.2%;
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1: FastSEQ for
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US-10-184-
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          DETECTION,
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Sequence 2342

Sequence 2342

Sequence 2342

Sequence 2342

Sequence 2342

Publication No. US20020198362A1

GENERAL INFORMATION:

APPLICANT: Gaiger, Alexander

APPLICANT: Algate, Paul A.

APPLICANT: Algate, Paul A.

APPLICANT: Mannion, Jane

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE DETECTION

TITLE OF INVENTION: HEMATOLOGICAL MALIGNANCIES

FILE REFERENCE: 2077. 001200

CURRENT APPLICATION NUMBER: US/09/796,692

CURRENT FILING DATE: 2001-03-01

PRIOR APPLICATION NUMBER: 60/190,479

PRIOR APPLICATION NUMBER: 60/200,545

PRIOR FILING DATE: 2000-04-27

PRIOR FILING DATE: 2000-04-28

PRIOR FILING DATE: 2000-04-28
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE DET
TITLE OF INVENTION: HEMATOLOGICAL MALIGNANCIES
FILE REFERENCE: 2077.001200
CURRENT APPLICATION NUMBER: US/09/796,692
CURRENT FILING DATE: 2001-03-01
PRIOR PILING DATE: 2000-03-17
PRIOR FILING DATE: 2000-03-17
PRIOR FILING DATE: 2000-03-17
PRIOR FILING DATE: 2000-04-27
PRIOR APPLICATION NUMBER: 60/200,303
PRIOR FILING DATE: 2000-04-28
PRIOR PELING DATE: 2000-04-28
PRIOR PELING DATE: 2000-04-28
PRIOR FILING DATE: 2000-04-28
PRIOR FILING DATE: 2000-04-28
PRIOR FILING DATE: 2000-05-01
PRIOR FILING DATE: 2000-05-02
PRIOR FILING DATE: 2000-05-02
PRIOR FILING DATE: 2000-05-02
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Mismatches
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PRIOR APPLICATION NUMBER: 60/223,378
PRIOR FILING DATE: 2000-08-07
NUMBER OF SEQ ID NOS: 9597
SOFTWARE: FastSEQ for Windows Version 3
SEQ ID NO 2277
LENGTH: 120
TYPE: PRT
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NAME/KEY: variant
LOCATION: (1)...(120)
OTHER INFORMATION: Xaa
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ORGANISM: Homo sapiens
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NUMBER: US/
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US-09-843-164-10
Sequence 10, Application US/09843164
GENERAL INFORMATION:
APPLICANT: Walke, D. Wade
APPLICANT: Wang, Xiaoming
TITLE OF INVENTION: NO. US20020061556Alel Human Membrane P
FILE REFERENCE: 07705.0014-00000
CURRENT APPLICATION NUMBER: US/09/843,164
CURRENT FILING DATE: 2001-04-27
PRIOR FILING DATE: 2000-04-27
NUMBER OF SEQ ID NOS: 19
SOFTWARE: FastSEQ for Windows Version 4.0
                                      GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
APPLICANT: Chen, Jian
APPLICANT: Chen, Jian
APPLICANT: Chen, Jian
APPLICANT: Goddard, Audrey
APPLICANT: Godowski, Paul J.
APPLICANT: Godowski, Paul J.
APPLICANT: Gurney, Austin L.
APPLICANT: Smith, Victoria
APPLICANT: Matanabe, Colin K.
APPLICANT: Watanabe, Colin K.
APPLICANT: Wood, William I.
APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPT]
TITLE OF INVENTION: ACIDS. ENCODING THE SAME
FILE REFERENCE: P3430R1C227
CURRENT APPLICATION NUMBER: US/10/184, 644
CURRENT FILING DATE: 2002-06-28
PLIOR APPLICATION TEMOVED - See File Wrapper or Palm
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Pred. No. 37;
5; Mismatches
Sequence 17, Application US/10184644
Publication No. US20030044930A1
GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 8, Application US/09843164
Patent No. US20020061556A1
GENERAL INFORMATION:
APPLICANT: Walke, D. Wade
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Best Local Similarity 55.6%;
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Best Local Similarity 45.0%;
Matches 9; Conservative
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NUMBER OF SEQ ID NOS: 612
SEQ ID NO 17
LENGTH: 1771
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US-10-184-644-17
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LENGTH: 423
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US20020061556Alel Human Membrane Proteins and Polynucleot
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Pred. No.
                                                                                                                                                                                                                                                                 60/199,950
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                                                                                                                  Application US/09843164
S20020061556A1
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                                                                                                                                                                                                                                                                                                SEQ ID NOS: 19
FastSEQ for Windows
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50.08;
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NTION: No. US200
E: 07705.0014-00
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9; Conservative
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Peter
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US-09-971-228-12
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Best Local S
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US-09-843-164-2
: Sequence 2, Application US/09843164
: APPLICANT: Walke, D. Wade
: APPLICANT: Wang, Xiaoming
: APPLICANT: Wang, Xiaoming
: TITLE OF INVENTION: NO. US20020061556Alel Human Membrane Pro
: FILE REFERENCE: 07705 0014-00000
: CURRENT APPLICATION NUMBER: US/09/843,164
: CURRENT FILING DATE: 2001-04-27
: PRIOR APPLICATION NUMBER: US 60/199,950
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US-09-843-164-12

Sequence 12, Application US/09843164

Patent NO. US20020061556A1

GENERAL INFORMATION:
APPLICANT: Wang, Xiaoming
APPLICANT: Wang, Xiaoming
APPLICANT: Scoville, John
TITLE OF INVENTION: NO. US20020061556A1el Human Mer
FILE REFERENCE: 07705 0014-00000
CURRENT APPLICATION NUMBER: US/09/843,164

CURRENT FILING DATE: 2001-04-27

PRIOR FILING DATE: 2000-04-27

NUMBER OF SEQ ID NOS: 19

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 12
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SOFTWARE: FastSEQ for Windows Version
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200 LSNVGCALSVTGLALTVIFQ 219
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Best Local Similarity 45.0%;
Matches 9; Conservative
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larity 45.0%;
Conservative
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NUMBER OF SEQ ID NOS: 81
SOFTWARE: Patentin version
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                         SEQ ID NO 2
LENGTH: 560
TYPE: PRT
ORGANISM: Homo sapiens
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US-09-843-164-12
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Sequence 2, Application US/09812272

Sequence 2, Application US/09812272

Sequence 1, O. US20020045214A1

GENERAL INFORMATION:

APPLICANT: Pamela Lane
APPLICANT
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US-10-037-616-27
Sequence 27, Application US/10037616
Patent No. US20020123148A1
GENERAL INFORMATION:
APPLICANT: English, Denis
APPLICANT: Rizzo, Maria.T.
APPLICANT: Rizzo, Maria.T.
APPLICANT: Sliva, Daniel T.
TITLE OF INVENTION: Sphingolipid Compositions and Methods fries Reference: 7042-119
CURRENT APPLICATION NUMBER: US/10/037,616
CURRENT FILING DATE: 2002-04-19
PRIOR PILING DATE: 2000-10-27
NUMBER OF SEQ ID NOS: 27
SOFTWARE: Patentin version 3.1
SEQ ID NO 27
LENGTH: 400
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Pred. No. 48;
3; Mismatches
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Job time : 5.82833 secs
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illarity 50.0%;
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; ORGANISM: Rattus norvegicus
US-10-037-616-27
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195 LFCVLAFLGILAAICA 210
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195 LFCVLAFLGILAAICA 210
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US-09-293-5C
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US-09-817-18
US-09-817-18
US-08-696-26
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1 NUMBER: US,
E: 2000-11-02
F: 09/116,641
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Pred.
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INVENTION: HUMAN ASPARTIC
RENCE: PF-0458-1 CIP
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N NUMBER: 09/008,
1: 1998-01-16
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LOCATION: 322
OTHER INFORMATION: :
\begin{smallmatrix} \mathcal{Q} & \mathcal{Q} 
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ORGANISM: HOMO :
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LENGTH: 433
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    /cgn2_6/ptodata/2/1aa/5B_COMB.pep:*
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    /cgn2_6/ptodata/2/1aa/FCTUS_COMB.pep:*
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US-08-974-691-2

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US-08-637-670-25

US-08-657-14

US-09-105-343A-2

US-09-105-3438-3

US-09-224-110-14

US-09-224-110-14

US-09-224-110-14

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US-09-224-110-14

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US-09-162-021B-2

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Jordan

APPLICANT

T: Tang, Jo
INVENTION:
F SEQUENCES:

Application US/08974691 6225103

Sequence

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Match Match 337.77 34.44 3

Length 433;

NUMBER: US 60/046,126 09-MAY-1997 INFORMATION:

DATA:

OMRF 166

NUMBER:

INFORMATION:

404-873-8795

FOR

31,284

MBER: US/08/974,691 20-NOV-1997

ICATION DATA:

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APPLICATION NUMBER: US 60/031,196 FILING DATE: 20-NOV-1996
                                                                    APPLICATION DATA:
                                                                                                                                         ICATION NUMBER:
                                                                                                         FILING DATE: 20
IOR APPLICATION
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                                                                                                                                                                         TORNEY/AGENT
                                                                                                                                                                                                          STRATION
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Patrea L. Pabst
STREET: 2800 One Atlantic Center, 1201 W. Peachtree
STREET: St.
CITY: Atlanta
STATE: GA
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. Patent No. 6225103
;    GENERAL INFORMATION:
    APPLICANT: Keolsch, Gerald
    APPLICANT: Lin, Xilli
    APPLICANT: Tang, Jordan
    TITLE OF INVENTION: Cloning and Characterization of
    NUMBER OF SEQUENCES: 14
    CORRESPONDENCE ADDRESS:
    ADDRESSEE: Patrea L. Pabst
    STREET: 2800 One Atlantic Center, 1201 W. D.C.
    STREET: St.
    CITY: Atlanta
    STATE: GA
                                                                                         COUNTRY: USA

2IP: 30309-3450

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/974,691

FILING DATE: 20-NOV-1997

CLASSIFICATION:
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5. 22;
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Pred. No. 2
3; Mismatch
                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 60/031,196
FILING DATE: 20-NOV-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/046,126
FILING DATE: 09-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Pabst, Patrea L.
REGISTRATION NUMBER: 31,284
REFERENCE/DOCKET NUMBER: OMRF 166
TELECOMMUNICATION INFORMATION:
TELEPHONE: 404-873-8794
TELEFAX: 404-873-8795
INFORMATION FOR SEQ ID NO: 6:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: USA
ZIP: 30309-3450
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
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Matches 9; Conservative
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LENGTH: 445 amino acir
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TOPOLOGY: 1
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INFORMATION FOR
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sequence 2, Ap
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V: Cloning and Characterization of Napsin
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2800 One Atlantic Center, 1201 W.
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09-MAY-1997
INFORMATION:
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ICATION NUMBER: US/08/974,691
NG DATE: 20-NOV-1997
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Release #1.0,
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Best Local Similarity 42.98;
Matches 9; Conservative
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CHARACTERISTICS
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READABLE
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OPERATING
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US-08-974-691-3
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FOR TREATMENT AND DIAGNOSIS
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ATE: 12-FEB-1998
GENT INFORMATION:
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Release #1.0,
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CORRESPONDENCE ADDRESS
ADDRESSEE: Abbott L
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                                                                                                                                                                                                                                                                          NUMBER:
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8; Conserv
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Best Local S
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US-08-637-670-25
; Sequence 25, Application US/08637670
; Patent No. 6413521
; GENERAL INFORMATION:
; APPLICANT: MCMICHAEL-PHILLIPS et al.
; TITLE OF INVENTION: Helminth Parasite Antigen with
; TITLE OF INVENTION: Aminopeptidase-like Activity
; WUMBER OF SEQUENCES: 43
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Barbara G. Ernst
; STREET: 555 13TH STREET, NW Suite 701E
; CITY: Washington
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US-09-705-448-10
Sequence 10, Application US/09705448
Patent No. 6432690
GENERAL INFORMATION:
APPLICANT: Xu, Hong
APPLICANT: Elsenboss, Laura A.
APPLICANT: Elsenboss, Laura A.
APPLICANT: Eogliano, Michael
APPLICANT: Cohan, Victoria L.
APPLICANT: Cohan, Victoria L.
APPLICANT: Bandman, Olga
TITLE OF INVENTION: HUMAN ASPARTIC PROTEASES
FILE REFERENCE: PF-0458-1 CIP
CURRENT APPLICATION NUMBER: US/09/705,448
CURRENT APPLICATION NUMBER: 09/116,641
PRIOR FILING DATE: 1998-07-16
PRIOR FILING DATE: 1998-01-16
NUMBER OF SEQ ID NOS: 10
SOFTWARE: FastSEQ for Windows Version 3:0
LENGTH: DATE
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Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                      Score 45;
REFERENCE/DOCKET NUMBER: OMRF 16
TELECOMMUNICATION INFORMATION:
TELEPHONE: 404-873-8794
TELEFAX: 404-873-8795
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 419 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; OTHER INFORMATION: 1906810, GenBank US-09-705-448-10
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Best Local Similarity 33.3%; Pumatches 7; Conservative 6;
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ORGANISM: MUS MUSCULUS
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                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 08/469,667 FILING DATE: 06-JUN-1995 TORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                          Release
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                                                                                              Becker Farm Road
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APPLICATION NUMBER: PC
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                                                                                                                                                                                                                                                                                                                                                                                                                           201-994-1744
                                                                                                                                                                                                                                                                                               DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
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                                                                                                                                                                     FORM:
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REGISTRATION NUMBER
REFERENCE/DOCKET NU
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                                                                                                                                                    ZIP: 07068-1739
COMPUTER READABLE DEDING TYPE: FLO
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GENERAL INFORMATION
APPLICANT: Yu, G
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                                                                                                               Roseland
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                                        NUMBER OF SEQUI
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                                                                ADDRESSEE:
ADDRESSEE:
STREET: 6
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US-09-224-110-14
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Best Local S
Matches 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/469,667
FILLING DATE: 06-JUN-1995
                                                                                                                            Score 43; DB
Pred. No. 33;
2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       36,134
R: 325800-435
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4: Colon Specific (
8: 24
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06-JUN-1995
                                                                                                                                                                                                                                                                              Sequence 14, Application US/08469667

Patent No. 5733748
GENERAL INFORMATION:
APPLICANT: Yu, Guo-Liang
APPLICANT: Rosen, Craig
TITLE OF INVENTION: Colon Specific
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
ADDRESSEE: Carella, Byrne, Bain, ADDRESSEE: Stewart & Olstein
STREET: 6 Becker Farm Road
CITY: Roseland
STATE: NJ
COUNTRY: USA
ZIP: 07068-1739
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Pred.
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TELEPHONE: 201-994-1700
; INFORMATION FOR SEQ ID NO: 2:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 249 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: No. 6207642e
US-09-105-343A-2
                                                                                     6207642e
                                                                                                                            35.2%;
^{\circ}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TTORNEY/AGENT INFORMATION:
NAME: FEITARO, GREGORY D
REGISTRATION NUMBER: 36,
REFERENCE/DOCKET NUMBER:
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clarity 52.9%;
Conservative
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CHARACTERISTICS:
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| ALCMLGLVLALLSSSSA 21
                                                                                                                                         10; Conservative
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Best Local Similarity
Matches 9; Conser
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CLASSIFICATION:
ATTORNEY/AGENT IN
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; MOLECULE TYPE:
US-08-469-667-14
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US-09-224-110-14
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US-08-469-667-14
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INFORMATION F
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Best Local 3
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Roger
TWO NOVEL HUMAN CATHESPIN PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US/09/080,538
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CATION NUMBER: 08/723,938
G DATE:
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ICATION INFORMATION:
E: 415-855-0555
415-845-4166
                                                      34.4%;
Similarity 38.1%;
8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION:
                                                                                                         QIHCALALLGLALAICSQGAA
                                                                                                                                 259 OTHMERVKVGPGLTLCAKGCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Billings, Lucy J
REGISTRATION NUMBER: 36,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OIHCALALLGLALAICSOGAA
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OPERATING SYSTEM: DO
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: LUNGNOT02
312099
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     ARE: FastSEQ
APPLICATION D
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8; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                               TYPE:
                                                                                                                                                                                                                                                                                                                                                                                 94304
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            ; CLONE:
US-08-723-938-3
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Best Local S
Matches 8
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; Sequence 3, Application US/08723938
; Patent No. 5776759
; GENERAL INFORMATION:
    APPLICANT: Bandman, Olga
    APPLICANT: Coleman, Roger
    TITLE OF INVENTION: TWO NOVEL HUMAN CATHESPIN PROTEINS
; VUMBER OF SEQUENCES: 4
; CORRESPONDENCE ADDRESS:
    ADDRESSEE: Incyte Pharmaceuticals, Inc.
    STREET: 3174 Porter Drive
    CITY: Palo Alto
    STATE: CA
                                                                                                                                                                                                                                      Ind
                                                                                                                                                                                                                  Length
                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Billings, Lucy J
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-0125 US
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
TELEFAX: 415-845-4166
                                                 5800-26
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t: US/08/723,938
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STATE: CA
COUNTRY: U.S.
ZIP: 94304
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ VERSION 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/723,9
FILING DATE:
CLASSIFICATION DATA:
APPLICATION NUMBER:
FILING DATE:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Ferraro, Gregory D.
REGISTRATION NUMBER: 36,134
REFERENCE/DOCKET NUMBER: 3258(
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-994-1744
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 81 amino acids
TYPE: amino acid
TOPOLOGY: linear
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TYPE: amino acid
STRANDEDNESS:
                                                                                                                                                                                                                34.4%;
Similarity 52.9%;
9; Conservative
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PCT-US95-07289-14
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US-08-723-938-3
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Length 420;
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                                                                                                                                                                                                                           ION NUMBER: US 60/031,196
ATE: 20-NOV-1996
                                                                                                                                                                                                                                                                 MBER: US 60/046,126
09-MAY-1997
                                                                                                                                                            MBER: US/08/974,691
20-NOV-1997
                                                                                                                        rsTEM: PC-DOS/MS-DOS
Patentin Release #1.0
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                                                                                                      compatible
PC-DOS/MS
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-07-16
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Cohan, Victoria L
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Best Local Similarity 38.1%;
Matches 8; Conservative
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                                                                                                                                                                                                                                                                                                                                                       NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   420 amino acids
ino acid
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                                                                                                                                       APPLICATION DATA:
APPLICATION NUMBER: US
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                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy d
COMPUTER: IBM PC COMP
OPERATING SYSTEM: PC-
SOFTWARE: Patentin Re
CURRENT APPLICATION DATA
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Elsenboss, Lau
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SEQUENCE CHARACTERISTIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               protein
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linear
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ORIGINAL SOURCE:
               Atlanta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    amino
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S-08-974-691-8
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FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                       TELECO
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Patent N
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APPLICANT: Lin, Xinli
APPLICANT: Tang, Jordan
TITLE OF INVENTION: Cloning and Characterization of Na
NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indel
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STREET: 2800 One Atlantic Center, 1201 W. Peachtree
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length
                                                                                                                                                                                                                                                                         STATE: CA
COUNTRY: USA
ZIP: 94304
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/008, 271A
FILING DATE: 16-Jan-1998
PRIOR APPLICATION DATA:
                                                                                                                                                                                  TITLE OF INVENTION: HUMAN PROTEASE MOLECULES NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals, Inc.
STREET: 3174 Porter Dr.
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Mohan-Peterson, Sheela
REGISTRATION NUMBER: 41,201
REFERENCE/DOCKET NUMBER: PF-0458 US
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-855-0555
TELEFAX: 650-845-4166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 42; DB
Pred. No. 85;
4; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   <Unknown>
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CLONE: 877617
SEQUENCE DESCRIPTION: SEQ ID NO:
US-09-008-271A-4
                                                                          Bandman, Olga
Hillman, Jennifer I
Yue, Henry
Guegler, Karl J.
Corley, Neil C.
Tang, Tom Y.
Shah, Purvi
                            Sequence 4, Application US/09008271A
Patent No. 6203979
GENERAL INFORMATION:
APPLICANT: Bandman, Olga
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 420 amino acids TYPE: amino acid
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US-08-974-691-8
; Sequence 8, Application US/08974691
; Patent No. 6225103
; GENERAL INFORMATION:
; APPLICANT: Keolsch, Gerald
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: <Unknown> ATTORNEY/AGENT INFORMATION:
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Shah,
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INFORMATION FOR SEQ
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RESULT 13
US-09-008-271A-4
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; OTHER INFORMATION: 372637, LUNGNOT02 US-09-705-448-1 Query Match
34.4%; Score 42; DB 4; Length 420;
Best Local Similarity 38.1%; Pred. No. 85;
Matches 8; Conservative 4; Mismatches 9; Indels 0; Gaps

Search completed: March 21, 2003, 11:42:36 Job time : 5.72103 secs

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Maximum Match 100%
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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glucagon; acquired epilepsy; Alzheimer neoplasm; metastasis anti-human immunodel
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to an Oncorhynchus mykiss somatostatin poly containing a portion of preprosomatostatin I (PPSS-I) and/or a preprosomatostatin I (PPSS-II). The protein sequences and their associated polynucleotides are useful for identifying medified comatostatin polypeptide which functions as a somatostatin agonis for research, therapeutics or diagnostics, including medical and veterinary applications. The wild-type somatostatin and its modification are useful for treating hypersecretion from endocrine tum the plunitary (e.g. acromegaly) or gastroenteropancreatic tissues gastrinoma, glucagonoma, carcinoid syndrome), to cause tumour shr through their effects on cell proliferation and apoptosis and as in the treatment of diabetes mellitus via inhibition of growth he associated with acquired immunodeficiency syndrome (AIDS) and verneurological disorders (e.g. epilepsy, Alzheimer's disease and Huntington's disease) and somatostatin antagonists are effective treatment of such conditions. Nucleic acids encoding the polypept cuseful in gene therapy and fusion peptides can be targeted to neouseful in gene therapy and fusion peptides can be targeted to neouseful in gene therapy and fusion peptides can be targeted to neouseful in gene therapy and fusion peptides can be targeted to neouseful associated with sequence represents 0. Mykiss PPSS-II'' protein.

This sequence represents 0. Mykiss PPSS-II' protein.
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Best Local Similarity 100.0%; Pred. No. 1e-10;
Matches 25; Conservative 0; Mismatches 0; Inde
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                                                                                                                                                                                                                                                                                                                                                                                                                                              New somatostatin polypeptides derived from Oncorhynchus for treating diabetes mellitus, acromegaly, gastrinoma, immunodeficiency syndrome and neurological disorders -
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/note= "Dibasic cleavage site"
98.111
/note= "SS-14 variant peptide"
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AAU07667 standard; Protein; 115 AA:
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04-DEC-2001 (first entry)
                                                                                   2000CA-2325169.
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B; AAS12935.
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AAU07667
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The invention relates to an Oncorhynchus mykiss somatostatin polypeptide containing a portion of preprosomatostatin I (PPSS-I) and/or a portion of preprosomatostatin II (PPSS-II). The protein sequences and their associated polynucleotides are useful for identifying modified somatostatin polypeptide which functions as a somatostatin agonist useful for research, therapeutics or diagnostics, including medical and veterinary applications. The wild-type somatostatin and its modified version are useful for treating hypersecretion from endocrine tumours in the pituitary (e.g. acromegaly) or gastroenteropancreatic tissues (e.g. gastrinoma, glucagonoma, carcinoid syndrome), to cause tumour shrinkage through their effects.
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-- "Prosomatostatin II'
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note= "Mature PPSS-II'"
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Pred. No.
                                                                                                                                                                                                                                                            'note= "Signal peptide"
cquired immunodeficiency
lzheimer's disease; Hunt
letastasis; gene therapy;
immunodeficiency virus;
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--- "PPSS-II' F
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N-PSDB; AAS12934.
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    dels
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s, acromegaly, gastrinoma,
neurological disorders -
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    H
                                                                                                                                                                                                                                                                                          trout; somatostatin; preprosomatostatin; hyp; PPSS-II'; endocrine tumour: pituitary alan
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   Mismatches
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1. 24
/note= "Signal peptide"
1. 88
/note= "PPSS-I pre-serman
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- "Dibasic cleavage
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   mote.
                                                                                                                                                                                                                         (first
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B; AAS12933.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ŝ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cleavage-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (NDSU-) NDSU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1; Fig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CA2325169-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              03-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sheridan MA,
                                                                                                                                                                                                                       04-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            03-JUN-2001
22;
                                                                                                                                                                                        AAU07666;
                                                                                                                                                                                                                                                                                             Rainbow t
PPSS-II';
                                                                                                                                                   AAU07666
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 200
N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Peptide
  Matches
                                                                                                                     RESULT 3
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che pitulitary (e.g. acromegaly) or gastroenteropancreatic tissues (e.g. gastrinoma, glucagonoma, carcinoid syndrome), to cause tumour shrinkage through their effects on cell proliferation and apoptosis and as adjuncts in the treatment of diabetes mellitus via inhibition of growth hormone and glucagon. In addition, dysfunctional somatostatin secretion is associated with acquired immunodeficiency syndrome (AIDS) and various neurological disorders (e.g. epilepsy, Alzheimer's disease and Huntington's disease) and somatostatin antagonists are effective in the treatment of such conditions. Nucleic acids encoding the polypeptides are useful in gene therapy and fusion peptides can be targeted to neoplasms and their metastases, inhibiting the release of their secretory products. This sequence represents O. Mykiss PPSS-I protein.

Note: The features for this sequence are specifically claimed in the specification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This sequence represents the wild type human enolase (NSE) protein. The primers represented by AAT60282-T60285 were used to amplify tyrosine introduced versions of this sequence. The introduced tyrosine residues combine a radioisotope in such a way as to have no substantial effect or the antigenicity of the protein. The peptides with the introduced tyrosine residue has been introduced via an insertion or substitution into the wild type sequence. The peptides can then he means in radioimmunoscent
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Introducing tyrosine residues into a protein for radioisotopic labelling - by substitution, addition and/or insertion to a DNA coding for the protein, antigenicity of the protein is unchanged by labelling
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              expressing a recombinant DNA seque
een introduced via an insertion or
sequence. The peptides can then be
roduction of the label does not all
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               human;
                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                                                                                                                                                                                                                                                                                                 Length 114;
                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                          core 64; DB 22;
red. No. 0.048;
Mismatches 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 type sequence. The introduction of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         polymerase chain reaction; primer; se; radioisotope; antigen.
                                                                                                                                                                                                                                                                                                                                                                                              Score 64;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               433 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MRVSQIHCALALLGLALAICSQGAA
| ::: |||||| ||||| || ||
| MLSTRVQCALALLSLALAISSVSAA
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                                                                                                                                                                                                                                                                                                                                                                                          52.5%;
larity 60.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    95JP-0145542
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              95JP-0145542
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        radioimmunoassays, as
antigenicity of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1997-059703/06.
                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
15; Conser
                                                                                                                                                                                                                                                                                                                                                114 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Enolase protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                JP08308584-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-MAY-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19-MAY-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-NOV-1996.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAW14001;
                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
Matches 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCR; poly enolase;
                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAW14001
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the
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AAW14001
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Sequence

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; gene therapy; vaccine; lymphoma
antigen; chronic lymphocytic leuk
n's lymphoma; non-Hodgkin's lymph
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sence of haematological malignancy relation the patient (an increased level Haematological malignant, is intion
                                                                                                                                 Human haematological malignancy-related antigen #1611
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                for the detection of hema c lymphocytic leukemia, l d T/B cell non-Hodgkin's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 48; DB
Pred. No. 12;
4; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            y related antigen. The methods of the presence of haematological obtained from the patient (an in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ein sequence
The methods o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hodgkin's · lymphoma,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1152; 1252pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      detection, diagnosis and therapy present sequence is the protein sequence in the protein sequence is the protein sequence in the protein sequence is the protein sequence in the protein sequence is sequenced.
                                    120
                                                                                                                                                                               malignancy; anti
homa; Hodgkin's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            39.3%;
ilarity 56.2%;
Conservative
                                                                                                                                                                 vascular;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                compared to an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cular lymphoma, Hodgkin
non-Hodgkin's lymphoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Compositions and methods for malignancies, e.g. chronic lymphoma and Hodgkin's and
                                                                                                    entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  invention are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  | |:||::|| ||
ANAILGVSLAVCKAGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Algate PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ALALLGLALAICSQGA
                                                                                                                                                                               haematological maligifollicular lymphoma;
                                                                                                    (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORIXA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-514842/56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               risk).
                                                                                                                                                                                                                                                               WO200164886-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Page
                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 polypeptide,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22-MAY-2000;
14-JUL-2000;
03-AUG-2000;
04-AUG-2000;
07-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                present
                                                                                                                                                                                                                                                                                                 07-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    icular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                   AAM81913;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ouery Match
Best Local S
Matches 9
                                   AAM81913
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAM81978
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sample
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (CORI-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaiger
                                                                                                                                                                 Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              follî
cell
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AAM81978
ID AAM8
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RESULT AAM8191
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                haplotypes
                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ţ,
    433;
                                                                                                                                                                                                                                                                                                               muscle); ENO3; single nucleotide
3; isogene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention provides the protein, cDNA and gea human enclase 3 (beta, muscle) isogene containing a nucleotide polymorphisms (SNPs). The sequences can be uthe haplotype of an individual and identify whether parare linked to certain diseases. The present sequence is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indel
                                    Indel
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     Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel genetic variants of enolase 3, (beta, muscle) g studying expression and function of the protein, and drugs to treat disorders of glycolytic pathway -
    DB 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 52; DB Pred. No. 12; 4; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB
    Score 52; DB Pred. No. 12; 4; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Parks KE;
                                                                                                                                                                                                                                                                                                                                                                                                                           'note= "optionally Ser"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "optionally Val"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                "optionally Pro"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "optionally Ile
                                                                                                                                                                                                                                                                               Human enolase 3(beta, muscle) protein.
                                                                                                                                                                                                                                                                                                                                                                                              Key Location/Qualifiers Misc-difference 71
                                                                                                                                                                               AAM48922 standard; Protein; 434 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 28; Fig 3; 90pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (GENA+) GENAISSANCE PHARM INC.
                                   4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Koshy B,
  42.6%;
Clarity 58:8%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-154721/20.
N-PSDB; AAK98531, AAK98532.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               42.6%;
llarity 58.8%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 02-JUL-2001; 2001WO-US20952
                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   | |:||::||:|
| 108 ANAILGVSLAVCKAGAA 124
                                                                                   | |:||::||:|
107 ANAILGVSLAVCKAGAA 123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                9 ALALLGLALAICSOGAA 25
                                                                  9 ALALLGLALAICSQGAA 25
                                                                                                                                                                                                                                                                                                                 Human; enolase 3(beta, mu
SNP; haplotype analysis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             'note-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                'note*
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10; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Misc-difference 332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Finkel K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                76
Query Match
Best Local Similarity
Matches: 10; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                434 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                             Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10-JAN-2002.
                                                                                                                                                                                                                                                                                                                                                               sapiens
                                                                                                                                                                                                                                                16-APR-2002
                                                                                                                                                                                                              AAM48922;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Duda A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                 Homo
                                                                                                                                                RESULT 5
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ntion comprise antigen(s) in

haematological malignancies. The uence of a human haematological hods of the present invention comp

to compositions

follicular

hematological la, lymphoma, f in's lymphoma -

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using

of

of the indicative

which can be treated us c leukaemia, lymphoma, Hodgkin's lymphoma and

non-Hodgkin's

Gaps

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Length 120;

22;

DB 12.

op

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ID NO:3656

SEO

(first

coagulation;

wound

burn;

Tue

9

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Human; open reading frame; ORFX; detection; cytostatic; hepatotropic; vulnerary; antipsoriatic; antiparkinsonian; nootropic; neuroprotective; anticonvulsant; osteopathic; antiarthritic; immunosuppressant; cardiant; immunostimulant; thrombolytic; coagulant; vasotropic; antidiabetic; hypotensive; dermatological; immunosuppressive; antiinflammatory; antityiral; antibacterial; antifungal; antirheumatic; antithyroid; antibacterial; antifungal; antirheumatic; antithyroid; antibacterial; antifungal; antirtheumatic; antithyroid; antionidegenerative disorder; osteoarthritis; graft vs host disease; neurodegenerative disorder; osteoarthritis; hypothyroidism; SCID; AIDS; cardiovascular disease; diabetes mellitus; hypothyroidism; SCID; AIDS; cholesterol ester storage; systemic lupus erythematosus; infection; severe combined immunodeficiency; malaria; autoimmune disorder; asthma;
                                                                                                                                                                                                                                                                                                                       allergy, aplastic anaemia; nocturnal haemoglobinuria; bu bone damage; cartilage damage; antiinflammatory disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel nucleic acids and peptides derived from or useful for treating e.g. cancers, proliferative neurodegenerative disorders and cardiovascular c
Human ORFX ORF1828 polypeptide seguence
                                                                                                                                                                                                                                                                                                                                                                       thrombosis; contraceptive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2000-602362/57.
N-PSDB; AAC76273.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200058473-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     31-MAR-1999;
02-APR-1999;
05-APR-1999;
30-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           31-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-OCT-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ntion comprise
antigen(s) in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              be treated using
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   lymphoma and T/B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     follicular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        of
                                                                                                                                              leukaemia;
lymphoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    malignancies. Tlandamatological sent invention co
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          indicative
                                                                                                                        ymphoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Compositions and methods for the detection of hematological malignancies, e.g. chronic lymphocytic leukemia, lymphoma, lymphoma and Hodgkin's and T/B cell non-Hodgkin's lymphoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ch 162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              thods
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               nia,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to compositions and methodetection, diagnosis and therapy of haematological malignessent sequence is the protein sequence of a human haemalignancy related antigen. The methods of the present detecting the presence of haematological malignancy related sample obtained from the patient (an increased level polypeptide, compared to an unaffected individual, is in increased risk). Haematological malignancies which can the present invention are chronic lymphocytic leukaemia, follicular lymphoma, Hodgkin's lymphoma, non-Hodgkin's lymphoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      level
                                                                                                              Human; cytostatic; vascular; gene therapy; vaccine;
haematological malignancy; antigen; chronic lymphocy
follicular lymphoma; Hodgkin's lymphoma; non-Hodgkin
                                                                    Human haematological malignancy-related antigen #167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Lengt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 22;
17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 48; DB; Pred. No. 17; 4; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Page 1178; 1252pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ٦
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2000US-0190479.
2000US-0200545.
2000US-0200779.
2000US-0200779.
2000US-0200999.
2000US-0218950.
2000US-0218950.
2000US-0218950.
2000US-0218950.
2000US-0218950.
                                                                                                                                                                                                                                                                                                                                                                 2001WO-US07272
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      39.3%;
larity 56.2%;
Conservative
                       entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A, . Algate PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-514842/56.
                  (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORIXA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
Matches 9; Conser
                                                                                                                                                                                                                                                                   WO200164886-A2.
                                                                                                                                                                                                                                                                                                                                                                                                        01-MAR-2000;
27-APR-2000;
28-APR-2000;
28-APR-2000;
01-MAY-2000;
04-MAY-2000;
22-MAY-2000;
22-MAY-2000;
04-MG-2000;
03-AUG-2000;
04-AUG-2000;
                                                                                                                                                                                                                    Homo saptens
                                                                                                                                                                                                                                                                                                                                                              01-MAR-2001;
               13-NOV-2001
                                                                                                                                                                                                                                                                                                                 07-SEP-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (CORI-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaiger
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99US-0127607. 99US-0127636. 99US-0127728. 2000US-0540763.

Leach M;

2000WO-US08621

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AAC7446 to AAC77606 encode the proteins given in AAB40237 to AAB43397, which represent the human ORFX open reading frames 1 to 3161. The ORFX sequences have activities such as: cytostatic; hepatotropic; vulnerary; antipsoriatic; anticonvulsant; antiarthritic; immunosuppressant; cardiant; cardiant; thrombolytic; coagulant; vasotropic; antidiabetic; hypotensive; dermatological; immunosuppressive; antithflammatory; antibacterial; antiviral; antifungal; antirheumatic; antithflammatory; antibacterial; antiviral; antifungal; antitheumatic; cantinflammatory; antibacterial; antiviral; antifungal; antitheumatic; cantitheoremic of or predisposition to, or preventing or treating the presence of or predisposition to, or preventing or treating pathological conditions associated with an ORFX-associated disorder. The nucleic acids can be used to express ORFX proteins in gene therapy vectors. The proteins and nucleic acids may be used to treat cancers, proliferative disorders, neurodegenerative disorders, osteoarthritis, graft vs host disease, cardiovascular disease, diabetes mellitus, hypothyroidism, cholesterol ester storage, systemic lupus
                                                                                                                                                                                                                                                                                                                                                        The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            damage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  viral,
asthma,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cartilage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AIDS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            itciency (SCID), AIDS
autcimmune disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 429;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            contraceptive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ion, malaria, autoimmune di
i, burns, wounds, bone and
antiinflammatory disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   erythematosus, severe combined immunodeficiency bacterial or fungal infection, malaria, autoimmu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ø
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 48; DB 2
Pred. No. 48;
4; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   inhibit thrombosis; and as
     English
   5507pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    tic anaemia,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          39.38;
56.28;
2810-2811;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  bacterial or fungal infectablergies, aplastic anaem nocturnal haemoglobinuria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            coagulation; to
   Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   429
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local S
Matches 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
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Gaps

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Ind

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124

ALALLGLALAICSQGA | |:||:||:| | ANAILGVSLAVCKAGA

109

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8

RESULT 8

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24

429

AAB42064 standard; Protein;

AAB42064

AX X X

open reading 1 ve disorders, r disease -

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Gaps

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Indels

4;

Conservative

Similarity 9; Conservation

AAW54357;

103

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57

RESULT AAW543

6

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us-09-727-739b-19.rag

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The invention relates to gene sequences (ABI99915-ABI99934) having modified expression in fetal heart tissue as compared to adult heart tissue and the encoded proteins (ABB57375-ABB57392). The genes have cardiant activity and may be useful in the promotion of the repair of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          screening potential compounds for the ability to inted with myocardial necrosis. Drugs identified by tods may be used to treat and prevent disease with whose is associated, such as cardiac hypertrophy and ciency. Diagnosis of diseases such as those above.
                                                                                                                                                                                                                                                                                      in fetal and adult heart
for promoting repair of d
                                                                                                                                                                                                                                                                                                                                                                                                                                                     tul in the promotion of myocardial necrosis m
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 48; DB 23;
Pred. No. 49;
4; Mismatches
                                                                                                                                                                                         Sakurada K;
                                                                                                                                                                                                                                                                                                                                                                   Japanese
                                                                                                                                                                                                                                                                                   Genes having differential expression useful for screening potential drugs caused by myocardial necrosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           210 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                pneumonia; b
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chlamydia pneumoniae protein not
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 caused by
                                                                                                                                                                                         ×
                                                                                                                                                                                         Kikuchi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Respiratory disease; pneumonic
sinusitis; purulent otitis med
vaccine; neutralising epitope
                                                                                                                                                                                                                                                                                                                                                                Claim 53; Page 98-100; 171pp;
                                                                         27-APR-2001; 2001WO-JP03700.
                                                                                                                                                (KYOW ) KYOWA HAKKO KOGYO KK
                                                                                                             2000JP-0126741.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAY35223 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             98WO-IB01890
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ch 39.3%;
1 Similarity 56.2%;
7 9; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98US-0107078
97FR-0014673
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ALALLGLALAICSOGA 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              damage to heart tissue
are useful for screenin
disease associated with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cardiac insufficiency disclosed.
                                                                                                                                                                                       Sekine S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chlamydia pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first
                                                                                                                                                                                                                          WPI; 2002-075160/10.
N-PSDB; ABI99919.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        434 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 W09927105-A2.
                                                                                                             27-APR-2000;
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21-NOV-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-NOV-1998;
                                     08-NOV-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       03-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Griffais R;
                                                                                                                                                                                         Yamada Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             screening
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches · / 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAY35223;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 11
AAY35223
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           dd
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cardiac hypertrophy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ide SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  433;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Proteins AAW54349-W54364 are examples of proteins produced endometrium during the hyperplasia, adenocarcinoma or proliphase of the endometrium. The presence and quantities of total be detected using 2D gel electrophoresis comparison of The proteins can be used as biochemical markers to detect the endometrium and can be measured in body fluids, obviation endometrial biopsies.
                                                                                                                                                                                                                                                                 Endometrium; hyperplasia; adenocarcinoma; proliferative 2D gel electrophoresis; detection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 48; DB 19; Length
Pred. No. 48;
; Mismatches 3; Inde
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Rat mucocardial cell proliferation associated polypept
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           for,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Biochemical markers of human endometrium - diagnosis of hyperplasia and adenocarcinoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         myocardial necrosis;
                                                                                                             433 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABB57379 standard; Protein; 434 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 21; 77pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLINICAL & BASIC RES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Larsen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               97GB-0007132.
96GB-0018600.
                                                                                                            protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ouery Match 39.3%;
Best Local Similarity 56.2%;
Aatches 9; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                           97WO-GB02394
                                                                                                                                                                                      (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                107 ANAILGVSLAVCKAGA 122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             9 ALALLGLALAICSOGA 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Rat; heart; cardiant; reardiac insufficiency.
| |:||::||:|
| ANAILGVSLAVCKAGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Fey SJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1998-207057/18.
                                                                                                            AAW54357 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          433 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rattus norvegius
                                                                                                                                                                                                                               Alpha Enolase
                                                                                                                                                                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                             WO9810291-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (CLIN-) CENT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               08-APR-1997;
06-SEP-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Byrjalsen I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          08-MAR-2002
                                                                                                                                                                                       14-AUG-1998
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Gaps

Indels

Length 434;

disease; sarcoidosis;); pharyngitis;

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ABB57379;

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RESULT 10 ABB57379

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Sequence

trachomatis

sequences o influence

The gene ability to

the repair

identified by the

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These NGPCR's are hea, testis, liver

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Indels

9

Score 47; DB 23; Pred. No. 67; ; Mismatches

5

Length 423;

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The invention relates to 1 of 232 isolated or recombinant polynucleotides encoding an Arabidopsis thaliana transcription factor, their variants, complements, fragments, or related polynucleotide with 31% to 95% sequence identity, where the plant possesses an altered trait as compared to a wild-type or reference plant, or the plant exhibits an altered to a wild-type or reference plant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      An isolated or recombinant polynucleotide used to
 in signal transduction after ligand binding expressed in human placenta, bone marrow, t kidney cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    chemical;
                                                                                                                                                                                                                                                                                                                                          AAU93172 standard; Protein; 472
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              M, Creelman R, Dubell
Ratcliff O, Reuber JL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                864-866; 941pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Agriculture; metabolic chemica
microbial disease resistance;
fruit yield; growth rate; leaf
plant; transcription factor; t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BIOTECHNOLOGY
                                                                                                                                                                                                  VSQIHCALALLGLALAICSQ.22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2000US-227439P.
2000US-0713994.
2001US-0837944.
                                                                                                                Similarity 45.0%;
9; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2001WO-US26189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 transcription
                                                                                                                                                                                                                        Arabidopsis thallana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MENDEL BIC.
PILGRIM M.
CREELMAN R.
                                                                                                                                                                                                                                                                                                                                                                                                                        (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2002-292022/33
B; ABK65358.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RATCLIFF O
                                                                               423 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       YU G.
PINEDA O.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HEARD J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 40; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    KEDDIE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200215675-A1
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16-NOV-2000;
16-APR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22-AUG-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DAM
                                                                                                                                                                                                                                                                                                                                                                                                                        02-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Arabidopsis
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                                                                                                                    Query Match
Best Local S
Matches 9
                                                                                                                                                                                                                                                                                                                                                                             AAU93172;
                                                                               Sequence
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(PILG/
(CREE/
(DUBE/
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(RIEC/
(YUGG/
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                                                                                                                                                                                                                                                                                                    RESULT 13
AAU93172
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SSSXS
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0
                                                                                                                                                                                                                                 The open reading (79) can be used in Ing C. pneumoniae compositions,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel isolated G-protein coupled receptor polypeptide useful for identifying compounds that modulate the protein expression or activity which are used to treat a variety of physiological or mental disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The sequences given in AAB47761-69 show novel G-protein coupled receptor (NGPCR) polypeptides. The NGPCR coding sequences are found chromosome 3 at 3q11.1 in the human genome. These NGPCR proteins sho the presence of 7 conserved transmembrane domains which are involved interconnected by non-conserved hydrophilic loops. They are involved
                                                                                                                                        open readi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            bone marrow;
                                                                                                                                                                                                                      erythema
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                                                                                                                                                                                                  in heart
                                                                                                                                                                                                                                                                                                                                                                                                                    Length 210;
                                                                                                                              AAY34584-Y35879 represent the proteins encoded by all the of frames in the complete genome (see AAX91990) of Chlamydia p C. pneumoniae causes respiratory disease such as pneumoniae bronchitis and is thought to be a contributing factor in he disease, sarcoidosis, sinusitis, purulent otitis media, ery nodosum or pharyngitis. The polypeptides encoded by the operframes of the C. pneumoniae genome (see AAY34584-Y35879) calmmunogenic compositions as vaccines. Vectors containing C. nucleotides sequences can also be used as immunogenic compoespecially where the vector directs the expression of a neuepitope of C. pneumoniae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel G-protein coupled receptor; NGPCR; transmembransignal transduction; ligand binding; human; placentatrachea; testis; liver; kidney cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           6;
                                                                                                                                                                                                                                                                                                                                                                                                                    Score 47; DB 20;
Pred. No. 32;
5; Mismatches
                                                                                            Page 1070-1071; Disclosure; 1912pp; English.
                                                      Genome sequence of Chlamydia pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 78-79; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel G-protein coupled receptor #5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAB47765 standard; Protein; 423 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Scoville J;
                                                                                                                                                                                                                                                                                                                                                                                                                    38.5%; S
Similarity 45.0%; P
9; Conservative 5;
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:| |:|:||| :| 1
48 EIASAIAILGLLVAFCASAA 67
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N-PSDB; AAI72010.
              WPI; 1999-357842/30.
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                                                                                                                                                                                                                                                                                                                                                                           210 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200181411-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
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Best Local :
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AAB47765
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senescence

stress; dr nce; seed

emical; environmental stres ace; herbicide resistance; leaf senescence; flower se ar; transgenic.

transgenic

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produce

English.

Keddie Pineda O

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AJ, hear Riechmann

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us-09-727

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yield,
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exhibits ectopic expression or altered expression of one o associated with a plant trait as compared to a wild plant. are a transgenic plant comprising the polynucleotides, a creadable medium having stored sequence information, and id homologue sequence from a database comprising a plurality sequences comprising inputting sequence information select 464 fully defined sequences given in the specification. The recombinant polynucleotide is used for producing a plant hamodified trait, the method comprising selecting a plant hamodified trait, the method comprising selecting a polynucleotide or antisense nucleic acid into an expression introducing the vector into a plant or a cell of a plant the polypeptide or antisense nucleic acid, thereby product plant, and selecting for a modified trait (e.g. increased production of agriculturally useful proteins or metabolic pest tolerance, environmental stress response (e.g. drough disease resistance, herbicide resistance, seed and fruit y rate, leaf and flower senescence and many other traits lis specification). The present sequence is one of the 232 pro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Agriculture; metabolic chemical; environmental stress; microbial disease resistance; herbicide resistance; see fruit yield; growth rate; leaf senescence; flower senes plant; transcription factor; transgenic.
                                                                                                                                                                                                                                                                                                                                     Length
                                                                                                                                                                                                                                                                                                                                    23;
                                                                                                                                                                                                                                                                                                                               Score 47; DB 2
Pred. No. 75;
7; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Arabidopsis transcription factor #56
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                47
                                                                                                                                                                                                                                                                                                                                 Juery Match
Jest Local Similarity 40.9%;
Matches 9; Conservative 7
                                                                                                                                                                                                                                                                                                                                                                                          4 SQIHCALALGLALAICSOGAA
:111 |::: |:| |: |1|
193 AQIHAAVSVAGVAAAVAAIAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAU93018 standard; Protein;
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16-NOV-2000; 2000US-0713994
16-APR-2001; 2001US-0837944
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22-AUG-2001; 2001WO-US26189
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(CREE/) CREELMAN R.
(DUBE/) DUBELL A J.
(HEAR/) HEARD J.
(JIAN/) JIANG C.
(KEDD/) KEDDIE J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REUBER J L
RIECHMANN
                                                                                                                                                                                                                                                                                                        472 AA;
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PINEDA O.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RATCLIFF
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                                                                                                                                                                                                                                                                                                        Sequence
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(RIEC/)
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Matches
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AAU93018
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polynucleotides
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included
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one of
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all of a plant to overexpress
thereby producing a modified
e.g. increased
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it), microbial
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! yield, growth
!sted in the
                                                                                                                                          transgenic
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                                                                                                                                                                                                                                                                                                                                                    plant
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                             Pineda
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                                                                                                                                                                                                                                                                                                                                                                                                                                          information, and intrinsing a plurality
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reference plant,
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me plant possesses an altere
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          Jiang
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             inputting sequence information quences given in the specificati eotide is used for producing a pmethod comprising selecting a poe or an antisense nucleic acid,
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          AJ, Heard
Riechmann
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on). The present sequence is one
transcription factors.
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                                                                                                                                       polynucleotide
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ligand binding; huma
r; kidney cell.
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Pred. No.
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a wild-typ
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            Dubel1
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                                                                                                                                                                                                                                                                                                                                                                        expression
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                                                                                                                                                                                                                                                                                                                                                                                            a plant
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                                                                                                                                                                                                     40; Page 265-267;
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                                                                                                                                                                                                                                                                 Arabidopsis
fragments,
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          Pilgrim M, Creelman
Adam L, Ratcliff O,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        selecting
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B; ABK65204.
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(LEXI-) LEXICON GENETICS INC.
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Scoville J; Walke DW, × Wang

WPI; 2002-049266/06 N-PSDB; AAI72009.

Novel isolated G-protein coupled receptor polypeptid identifying compounds that modulate the protein exprwhich are used to treat a variety of physiological o

Claim 6; Page 77; 85pp; English.

The sequences given in AAB47761-69 show novel G-protein receptor (NGPCR) polypeptides. The NGPCR coding sequence chromosome 3 at 3q11.1 in the human genome. These NGPCR the presence of 7 conserved transmembrane domains which interconnected by non-conserved hydrophilic loops. They in signal transduction after ligand binding. These NGPC expressed in human placenta, bone marrow, trachea, testikidney cells.

Length 481; 6; Indels Score 47; DB 23; Pred. No. 76; 5; Mismatches 6 Query Match
Best Local Similarity 45.0%;
Matches 9; Conservative

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2003, 11:38:04 Search completed: March 21, Job time: 17.4506 secs 7

481 AA; Sequence

200 LSNVGCALSVTGLALTVIFQ 219 VSQIHCALALLGLALAICSQ 22 m